

**Table 3P Genes Corresponding to Hypertension Only 577**

Spot	p-value	Description	Accession	Unigene
55	0.023066	histone deacetylase 3 (HDAC3), mRNA /cds=(56,1342) /gb=NM_003883 /gi=13128861 /ug=Hs.446552 /len=1955	NM_003883	Hs.446552
56	0.033276	heterogeneous nuclear ribonucleoprotein D-like (HNRPDL), transcript variant 1, mRNA /cds=(581,1843) /gb=NM_005463 /gi=14110410 /ug=Hs.372673 /len=3514	NM_005463	Hs.372673
139	0.019991	fusion, derived from t(12;16) malignant liposarcoma (FUS), mRNA /cds=(79,1659) /gb=NM_004960 /gi=4826733 /ug=Hs.99969 /len=1824	NM_004960	Hs.99969
161	0.033276	putative nucleic acid binding protein RY-1 (RY1), mRNA /cds=(28,495) /gb=NM_006857 /gi=24307918 /ug=Hs.54649 /len=1433	NM_006857	Hs.54649
162	0.002441	mitogen-activated protein kinase kinase kinase 7 interacting protein 2 (MAP3K7IP2), transcript variant 2, mRNA /cds=(176,1786) /gb=NM_145342 /gi=21735558 /ug=Hs.109727 /len=4359	NM_145342	Hs.109727
188	0.034783	hypothetical protein FLJ25270 (FLJ25270), mRNA /cds=(244,1353) /gb=NM_152520 /gi=22749086 /ug=Hs.6295 /len=2493	NM_152520	Hs.6295
290	0.01344	Niemann-Pick disease, type C2 (NPC2), mRNA /cds=(116,571) /gb=NM_006432 /gi=20149580 /ug=Hs.433222 /len=929	NM_006432	Hs.433222
301	0.029076	chaperonin containing TCP1, subunit 4 (delta) (CCT4), mRNA /cds=(1,1620) /gb=NM_006430 /gi=5453604 /ug=Hs.79150 /len=1883	NM_006430	Hs.79150
325	0.036347	thyroid autoantigen 70kDa (Ku antigen) (G22P1), mRNA /cds=(656,2485) /gb=NM_001469 /gi=20070134 /ug=Hs.197345 /len=2743	NM_001469	Hs.197345
381	0.027779	clusterin (complement lysis inhibitor, SP-40,40, sulfated glycoprotein 2, testosterone-repressed prostate message 2, apolipoprotein J) (CLU), mRNA /cds=(48,1397) /gb=NM_001831 /gi=4502904 /ug=Hs.75106 /len=1676	NM_001831	Hs.75106
382	0.027779	accessory protein BAP31 (DXS1357E), mRNA /cds=(137,877) /gb=NM_005745 /gi=10047078 /ug=Hs.291904 /len=1314	NM_005745	Hs.291904
490	0.010364	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 2, 8kDa (NDUFA2), mRNA /cds=(57,356) /gb=NM_002488 /gi=4505354 /ug=Hs.163867 /len=590	NM_002488	Hs.163867
538	0.005658	choline kinase-like (CHKL), transcript variant 2, mRNA /cds=(185,568) /gb=NM_152253 /gi=23238260 /ug=Hs.154886 /len=4914	NM_152253	Hs.154886
665	0.033276	procollagen C-endopeptidase enhancer 2 (PCOLCE2), mRNA /cds=(197,1444) /gb=NM_013363 /gi=16904386 /ug=Hs.8944 /len=1988	NM_013363	Hs.8944

842	0.047	sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3E (SEMA3E), mRNA /cds=(467,2794) /gb=NM_012431 /gi=6912649 /ug=Hs.212414 /len=6474	NM_012431	Hs.212414
903	0.011512	S100 calcium binding protein A10 (annexin II ligand, calpactin I, light polypeptide (p11)) (S100A10), mRNA /cds=(112,405) /gb=NM_002966 /gi=4506760 /ug=Hs.400250 /len=649	NM_002966	Hs.400250
926	0.049002	similar to ribosomal protein L8 [Mus musculus] (LOC346158), mRNA	XM_294080	
950	0.036347	RAP1B, member of RAS oncogene family (RAP1B), mRNA /cds=(149,703) /gb=NM_015646 /gi=7661677 /ug=Hs.156764 /len=1927	NM_015646	Hs.156764
958	0.026531	cathepsin B (CTSB), transcript variant 2, mRNA /cds=(314,1333) /gb=NM_147780 /gi=22538430 /ug=Hs.297939 /len=2140	NM_147780	Hs.297939
1209	0.011512	membrane protein, palmitoylated 6 (MAGUK p55 subfamily member 6) (MPP6), mRNA /cds=(300,1922) /gb=NM_016447 /gi=21361597 /ug=Hs.108931 /len=2201	NM_016447	Hs.108931
1309	0.037968	oxysterol binding protein-like 11 (OSBPL11), mRNA /cds=(306,2549) /gb=NM_022776 /gi=23111058 /ug=Hs.61260 /len=4206	NM_022776	Hs.61260
1557	0.008362	MR2-CI0186-291100-010-a06 CI0186 cDNA, mRNA sequence /gb=BF814502 /gi=12147047 /ug=Hs.446594 /len=530	BF814502	Hs.446594
1573	0.037968	fibromodulin (FMOD), mRNA /cds=(21,1151) /gb=NM_002023 /gi=5016093 /ug=Hs.230 /len=2863	NM_002023	Hs.230
1746	0.023066	HSPC230 gene (HSPC230), mRNA /cds=(69,791) /gb=NM_016487 /gi=24475976 /ug=Hs.36069 /len=923	NM_016487	Hs.36069
1754	0.026531	hypothetical protein MGC13007 (MGC13007), mRNA /cds=(1099,1653) /gb=NM_032320 /gi=14150091 /ug=Hs.332382 /len=2479	NM_032320	Hs.332382
1758	0.011512	cyclin-dependent kinase inhibitor 1C (p57, Kip2) (CDKN1C), mRNA /cds=(261,1211) /gb=NM_000076 /gi=4557440 /ug=Hs.106070 /len=1511	NM_000076	Hs.106070
1786	0.049002	lamin A/C (LMNA), transcript variant 1, mRNA /cds=(213,2207) /gb=NM_170707 /gi=27436945 /ug=Hs.377973 /len=3181	NM_170707	Hs.377973
1829	0.036347	excision repair cross-complementing rodent repair deficiency, complementation group 5 (xeroderma pigmentosum, complementation group G (Cockayne syndrome)) (ERCC5), mRNA /cds=(198,3758) /gb=NM_000123 /gi=4503600 /ug=Hs.48576 /len=3854	NM_000123	Hs.48576
1854	0.026531	clone IMAGE:3611719, mRNA, partial cds /cds=(1,2592) /gb=BC003542 /gi=13097656 /ug=Hs.244482 /len=3234	BC003542	Hs.244482
1895	0.008591	kinesin family member 11 (KIF11), mRNA /cds=(141,3311) /gb=NM_004523 /gi=13699823 /ug=Hs.8878 /len=4908	NM_004523	Hs.8878

2098	0.011512	ubiquitin specific protease 13 (isopeptidase T-3) (USP13), mRNA /cds=(95,2686) /gb=NM_003940 /gi=4507848 /ug=Hs.85482 /len=2738	NM_003940	Hs.85482
2137	0.029076	NAD(P)H dehydrogenase, quinone 1 (NQO1), mRNA /cds=(51,875) /gb=NM_000903 /gi=4505414 /ug=Hs.406515 /len=2447	NM_000903	Hs.406515
2166	0.045065	nephroblastoma overexpressed gene (NOV), mRNA /cds=(73,1146) /gb=NM_002514 /gi=19923725 /ug=Hs.235935 /len=2389	NM_002514	Hs.235935
2206	0.00136	glycoprotein (transmembrane) nmb (GPNMB), mRNA /cds=(92,1774) /gb=NM_002510 /gi=4505404 /ug=Hs.82226 /len=2669	NM_002510	Hs.82226
2214	0.043196	ts79a05.x1 NCI_CGAP_GC6 cDNA clone IMAGE:2237456 3', mRNA sequence /clone=IMAGE:2237456 /clone_end=3' /gb=AI917390 /gi=5637245 /ug=Hs.99415 /len=462	AI917390	Hs.99415
2230	0.026531	PMS1 postmeiotic segregation increased 1 (S. cerevisiae) (PMS1), mRNA /cds=(81,2879) /gb=NM_000534 /gi=11496979 /ug=Hs.111749 /len=3121	NM_000534	Hs.111749
2248	0.008828	KIAA0155 gene product (KIAA0155), mRNA /cds=(87,3608) /gb=NM_014633 /gi=7661949 /ug=Hs.173288 /len=4243	NM_014633	Hs.173288
2254	0.034783	mRNA; cDNA DKFZp761F0118 (from clone DKFZp761F0118) /cds=(1,6490) /gb=AL831917 /gi=21732430 /ug=Hs.6685 /len=7334	AL831917	Hs.6685
2288	0.034783	Similar to hypothetical protein MGC30540, clone MGC:17342 IMAGE:4342258, mRNA, complete cds /cds=(216,1457) /gb=BC042899 /gi=27552863 /ug=Hs.153716 /len=3028	BC042899	Hs.153716
2326	0.039649	formin binding protein 4 (FNBP4), mRNA /cds=(28,3075) /gb=NM_015308 /gi=24308032 /ug=Hs.6834 /len=3995	NM_015308	Hs.6834
2329	0.004758	inhibitor of growth family, member 1-like (ING1L), mRNA /cds=(92,934) /gb=NM_001564 /gi=4504694 /ug=Hs.107153 /len=1078	NM_001564	Hs.107153
2408	0.037968	BCL2-associated athanogene 3 (BAG3), mRNA /cds=(307,2034) /gb=NM_004281 /gi=14043023 /ug=Hs.15259 /len=2605	NM_004281	Hs.15259
2437	0.039649	nm53h08.s1 NCI_CGAP_Br2 cDNA clone IMAGE:1071999 3' similar to gb:M10036 TRIOSEPHOSPHATE ISOMERASE mRNA sequence /clone=IMAGE:1071999 /clone_end=3' /gb=AA573489 /gi=2348017 /ug=Hs.404623 /len=230	AA573489	Hs.404623
2478	0.043196	ECSIT (LOC51295), mRNA /cds=(78,1373) /gb=NM_016581 /gi=20149632 /ug=Hs.22199 /len=1668	NM_016581	Hs.22199
2501	0.036347	replication protein A2, 32kDa (RPA2), mRNA /cds=(292,1104) /gb=NM_002946 /gi=21314636 /ug=Hs.79411 /len=1750	NM_002946	Hs.79411
2563	0.029076	clone MGC:13446 IMAGE:4275731, mRNA, complete cds	BC009777	Hs.143198

2589	0.030424	ribosomal protein L39 (RPL39), mRNA /cds=(68,223) /gb=NM_001000 /gi=16306563 /ug=Hs.300141 /len=401	NM_001000	Hs.300141
2698	0.01344	topoisomerase (DNA) II alpha 170kDa (TOP2A), mRNA /cds=(127,4722) /gb=NM_001067 /gi=19913405 /ug=Hs.156346 /len=5698	NM_001067	Hs.156346
2716	0.006339	general transcription factor IIIC, polypeptide 3, 102kDa (GTF3C3), mRNA /cds=(94,2754) /gb=NM_012086 /gi=6912397 /ug=Hs.90847 /len=2961	NM_012086	Hs.90847
2773	0.049002	nucleosome assembly protein 1-like 1 (NAP1L1), transcript variant 1, mRNA /cds=(125,1300) /gb=NM_139207 /gi=21327707 /ug=Hs.302649 /len=3582	NM_139207	Hs.302649
2787	0.037968	meningioma expressed antigen 6 (coiled-coil proline-rich) (MGEA6), mRNA /cds=(315,2729) /gb=NM_005930 /gi=5174560 /ug=Hs.117242 /len=3676	NM_005930	Hs.117242
2822	0.047	Rho-related BTB domain containing 3 (RHOBTB3), mRNA /cds=(336,2171) /gb=NM_014899 /gi=7662355 /ug=Hs.10432 /len=4099	NM_014899	Hs.10432
2829	0.045065	mitochondrion, complete genome	NC_001807	
2836	7.42E-05	lactate dehydrogenase B (LDHB), mRNA /cds=(103,1107) /gb=NM_002300 /gi=22726178 /ug=Hs.234489 /len=1336	NM_002300	Hs.234489
2880	0.031823	ribosomal protein L36A (RPL36A), mRNA /cds=(41,361) /gb=NM_021029 /gi=16306564 /ug=Hs.178391 /len=425	NM_021029	Hs.178391
2946	0.047	CD164 antigen, sialomucin (CD164), mRNA /cds=(94,687) /gb=NM_006016 /gi=21361273 /ug=Hs.43910 /len=3038	NM_006016	Hs.43910
2972	0.034783	cDNA FLJ39487 fis, clone PROST2015246. /gb=AK096806 /gi=21756379 /ug=Hs.273138 /len=2419	AK096806	Hs.273138
3013	0.049002	uncharacterized hematopoietic stem/progenitor cells protein MDS031 (MDS031), mRNA /cds=(35,532) /gb=NM_018466 /gi=20070304 /ug=Hs.110853 /len=1358	NM_018466	Hs.110853
3019	0.047	hypothetical protein FLJ22332 (FLJ22332), mRNA /cds=(617,1597) /gb=NM_024724 /gi=20127632 /ug=Hs.111092 /len=1719	NM_024724	Hs.111092
3039	0.010925	paternally expressed 10 (PEG10), mRNA /cds=(118,1095) /gb=NM_015068 /gi=14149662 /ug=Hs.137476 /len=6253	NM_015068	Hs.137476
3041	0.039649	replication factor C (activator 1) 5, 36.5kDa (RFC5), mRNA /cds=(102,1124) /gb=NM_007370 /gi=19923788 /ug=Hs.171075 /len=2097	NM_007370	Hs.171075
3048	0.037968	retinal short-chain dehydrogenase/reductase 2 (RetSDR2), mRNA /cds=(189,1091) /gb=NM_016245 /gi=7705904 /ug=Hs.12150 /len=1760	NM_016245	Hs.12150

3119	0.010925	heterogeneous nuclear ribonucleoprotein D-like (HNRPDL), transcript variant 1, mRNA /cds=(581,1843) /gb=NM_005463 /gi=14110410 /ug=Hs.372673 /len=3514	NM_005463	Hs.372673
3135	0.018141	mRNA; cDNA DKFZp586C1019 (from clone DKFZp586C1019) /gb=AL049397 /gi=4500188 /ug=Hs.12314 /len=1737	AL049397	Hs.12314
3171	0.036347	transforming, acidic coiled-coil containing protein 1 (TACC1), mRNA /cds=(321,2738) /gb=NM_006283 /gi=5454099 /ug=Hs.173159 /len=7758	NM_006283	Hs.173159
3232	0.00136	phosphoglycerate dehydrogenase (PHGDH), mRNA /cds=(137,1738) /gb=NM_006623 /gi=23308576 /ug=Hs.3343 /len=1968	NM_006623	Hs.3343
3248	0.029076	CDw92 antigen (CDW92), mRNA /cds=(43,2016) /gb=NM_080546 /gi=18034691 /ug=Hs.179902 /len=4301	NM_080546	Hs.179902
3262	0.00423	serine (or cysteine) proteinase inhibitor, clade F (alpha-2 antiplasmin, pigment epithelium derived factor), member 1 (SERPINF1), mRNA /cds=(39,1082) /gb=NM_002615 /gi=4505708 /ug=Hs.173594 /len=1199	NM_002615	Hs.173594
3275	0.041391	KIAA0141 gene product (KIAA0141), mRNA /cds=(81,1628) /gb=NM_014773 /gi=7661939 /ug=Hs.63510 /len=3020	NM_014773	Hs.63510
3287	0.007917	FtsJ 1 (E. coli) (FTSJ1), mRNA /cds=(301,1290) /gb=NM_012280 /gi=7110660 /ug=Hs.23170 /len=1867	NM_012280	Hs.23170
3347	0.036347	matrix metalloproteinase 13 (collagenase 3) (MMP13), mRNA /cds=(29,1444) /gb=NM_002427 /gi=13027796 /ug=Hs.2936 /len=2722	NM_002427	Hs.2936
3348	0.033276	family with sequence similarity 8, member A1 (FAM8A1), mRNA /cds=(56,1297) /gb=NM_016255 /gi=7705267 /ug=Hs.95260 /len=4695	NM_016255	Hs.95260
3370	0.002767	annexin A2 (ANXA2), mRNA /cds=(50,1069) /gb=NM_004039 /gi=4757755 /ug=Hs.217493 /len=1362	NM_004039	Hs.217493
3379	0.034783	jerky (mouse) (JRKL), mRNA /cds=(1,1575) /gb=NM_003772 /gi=22547223 /ug=Hs.105940 /len=2930	NM_003772	Hs.105940
3395	0.006705	mRNA for KIAA1423 protein, partial cds. /cds=(1,1852) /gb=AB037844 /gi=7243226 /ug=Hs.274396 /len=5390	AB037844	Hs.274396
3447	0.039649	duodenal cytochrome b (FLJ23462), mRNA /cds=(74,934) /gb=NM_024843 /gi=19923602 /ug=Hs.31297 /len=4254	NM_024843	Hs.31297
3464	0.019047	matrilin 3 (MATN3) precursor, mRNA /cds=(64,1524) /gb=NM_002381 /gi=13518040 /ug=Hs.278461 /len=2599	NM_002381	Hs.278461
3490	0.037968	FYN binding protein mRNA, complete cds	AF001862	Hs.58435

3508	0.047	hypothetical protein DKFZp762O076 (DKFZp762O076), mRNA /cds=(77,850) /gb=NM_018710 /gi=24308164 /ug=Hs.21621 /len=2266	NM_018710	Hs.21621
3516	0.045065	PM1-DT0054-231299-002-a09 DT0054 cDNA, mRNA sequence /gb=AW364737 /gi=6869491 /ug=Hs.407368 /len=643	AW364737	Hs.407368
3602	0.045065	thioredoxin-like 2 (TXNL2), mRNA /cds=(5,1012) /gb=NM_006541 /gi=5730103 /ug=Hs.42644 /len=1942	NM_006541	Hs.42644
3621	0.045065	tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, eta polypeptide (YWHAH), mRNA /cds=(198,938) /gb=NM_003405 /gi=21464102 /ug=Hs.349530 /len=1775	NM_003405	Hs.349530
3636	0.017272	paired immunoglobulin-like receptor beta (PILR(BETA)), mRNA /cds=(289,972) /gb=NM_013440 /gi=7305386 /ug=Hs.349256 /len=993	NM_013440	Hs.349256
3692	0.02533	cDNA FLJ13571 fis, clone PLACE1008405. /gb=AK023633 /gi=10435617 /ug=Hs.116278 /len=2484	AK023633	Hs.116278
3718	0.036347	dehydrogenase/reductase (SDR family) member 4 (DHRS4), mRNA /cds=(80,862) /gb=NM_021004 /gi=10337604 /ug=Hs.418501 /len=1281	NM_021004	Hs.418501
3759	0.027779	integrin-linked kinase (ILK), mRNA /cds=(157,1515) /gb=NM_004517 /gi=4758605 /ug=Hs.6196 /len=1789	NM_004517	Hs.6196
3766	0.019991	mRNA; cDNA DKFZp667G0425 (from clone DKFZp667G0425) /gb=AL832847 /gi=21733431 /ug=Hs.82254 /len=1961	AL832847	Hs.82254
3767	0.021999	cellular retinoic acid binding protein 2 (CRABP2), mRNA /cds=(138,554) /gb=NM_001878 /gi=6382069 /ug=Hs.183650 /len=969	NM_001878	Hs.183650
3796	0.02533	immunoglobulin lambda locus (IGL@) on chromosome 22	NG_000002	
3846	0.008362	myosin light chain 1 slow a (MLC1SA), mRNA /cds=(48,674) /gb=NM_002475 /gi=17986280 /ug=Hs.90318 /len=778	NM_002475	Hs.90318
3852	0.001188	matrilin 3 (MATN3) precursor, mRNA /cds=(64,1524) /gb=NM_002381 /gi=13518040 /ug=Hs.278461 /len=2599	NM_002381	Hs.278461
3857	0.045065	cDNA FLJ33181 fis, clone ADRGL2003684, highly similar to HLA CLASS I HISTOCOMPATIBILITY ANTIGEN, ALPHA CHAIN H PRECURSOR. /gb=AK090500 /gi=21748675 /ug=Hs.379218 /len=2290	AK090500	Hs.379218
3925	0.024176	similar to RIKEN cDNA 1110059E24, clone IMAGE:5218126, mRNA /gb=BC028019 /gi=20380167 /ug=Hs.112993 /len=3343	BC028019	Hs.112993
3947	0.045065	lung type-I cell membrane-associated glycoprotein (T1A-2), mRNA /cds=(234,722) /gb=NM_006474 /gi=18767663 /ug=Hs.135150 /len=1081	NM_006474	Hs.135150

3948	0.027779	chromosome 15 clone RP11-263119 map 15q15, complete sequence	AC021754	
3983	0.043196	CDP-diacylglycerol synthase (phosphatidate cytidyltransferase) 2 (CDS2), mRNA /cds=(258,1595) /gb=NM_003818 /gi=22035625 /ug=Hs.24812 /len=2711	NM_003818	Hs.24812
3992	0.003986	Gene 33/Mig-6 (MIG-6), mRNA /cds=(213,1601) /gb=NM_018948 /gi=21314673 /ug=Hs.11169 /len=3099	NM_018948	Hs.11169
3994	0.01344	mitochondrial ribosomal protein L42 (MRPL42), transcript variant 3, nuclear gene encoding mitochondrial protein, mRNA /cds=(179,607) /gb=NM_172178 /gi=26667173 /ug=Hs.112110 /len=2093	NM_172178	Hs.112110
3999	0.018141	cDNA FLJ30843 fis, clone FEBRA2002574. /gb=AK055405 /gi=16550125 /ug=Hs.350872 /len=2780	AK055405	Hs.350872
4008	0.003535	jagged 1 (Alagille syndrome) (JAG1), mRNA /cds=(414,4070) /gb=NM_000214 /gi=4557678 /ug=Hs.91143 /len=5896	NM_000214	Hs.91143
4021	0.023066	transketolase-like 1 (TKTL1), mRNA /cds=(121,1794) /gb=NM_012253 /gi=7110726 /ug=Hs.102866 /len=2455	NM_012253	Hs.102866
4032	0.001891	DNA sequence from clone RP1-68D18 on chromosome 11p12-13 Contains ESTs, STSs and GSSs. Contains part of the SLC1A2 gene for three isoforms of solute carrier family 1 (glial high affinity glutamate transporter) member 2 and part of the CD44 gene for CD44 antigen (homing function and Indian blood group system) with five isoforms, complete sequence	AL133330	
4034	0.021999	nucleostemin (NS), mRNA /cds=(31,1680) /gb=NM_014366 /gi=26892284 /ug=Hs.279923 /len=1916	NM_014366	Hs.279923
4055	0.008828	baculoviral IAP repeat-containing 2 (BIRC2), mRNA /cds=(1160,3016) /gb=NM_001166 /gi=10880127 /ug=Hs.289107 /len=3496	NM_001166	Hs.289107
4064	0.023066	ubiquitin-like 5 (UBL5), mRNA /cds=(66,287) /gb=NM_024292 /gi=13236509 /ug=Hs.13836 /len=413	NM_024292	Hs.13836
4086	0.00709	ceruloplasmin (ferroxidase) (CP), mRNA /cds=(1,3198) /gb=NM_000096 /gi=4557484 /ug=Hs.296634 /len=3321	NM_000096	Hs.296634
4101	0.008362	SEC14-like 1 (S. cerevisiae) (SEC14L1), mRNA /cds=(304,2451) /gb=NM_003003 /gi=4506866 /ug=Hs.75232 /len=5434	NM_003003	Hs.75232
4119	0.014875	genomic DNA, chromosome 11q, clone:CMB9-103D18, complete sequences	AP000857	
4121	0.007493	no match		

4122	0.00423	serologically defined colon cancer antigen 10 (SDCCAG10), mRNA /cds=(482,1600) /gb=NM_005869 /gi=5031958 /ug=Hs.23557 /len=1857	NM_005869	Hs.23557
4124	0.024176	LOC286218 (LOC286218), mRNA	XM_212235	
4157	0.039649	Siah-interacting protein (SIP), mRNA /cds=(118,804) /gb=NM_014412 /gi=7656951 /ug=Hs.27258 /len=2435	NM_014412	Hs.27258
4162	0.031823	acid phosphatase 1, soluble (ACP1), transcript variant a, mRNA /cds=(776,1252) /gb=NM_004300 /gi=4757713 /ug=Hs.75393 /len=2222	NM_004300	Hs.75393
4192	0.037968	no match		
4202	0.049002	serum/glucocorticoid regulated kinase-like (SGKL), transcript variant 1, mRNA /cds=(416,1705) /gb=NM_013257 /gi=25168264 /ug=Hs.380877 /len=4155	NM_013257	Hs.380877
4261	0.009828	epithelial membrane protein 3 (EMP3), mRNA /cds=(242,733) /gb=NM_001425 /gi=4503562 /ug=Hs.9999 /len=817	NM_001425	Hs.9999
4263	0.014875	testis expressed gene 292 (FLJ14728), mRNA /cds=(49,2109) /gb=NM_032830 /gi=14249535 /ug=Hs.151001 /len=2192	NM_032830	Hs.151001
4298	0.041391	pannexin 1 (PANX1), mRNA /cds=(349,1617) /gb=NM_015368 /gi=7662507 /ug=Hs.30985 /len=2738	NM_015368	Hs.30985
4314	0.020975	protein phosphatase 4, regulatory subunit 1 (PPP4R1), mRNA /cds=(94,2895) /gb=NM_005134 /gi=4826933 /ug=Hs.3382 /len=3878	NM_005134	Hs.3382
4319	0.017272	aryl hydrocarbon receptor interacting protein (AIP), mRNA /cds=(118,1110) /gb=NM_003977 /gi=4502008 /ug=Hs.75305 /len=1244	NM_003977	Hs.75305
4343	0.012127	lectin, galactoside-binding, soluble, 1 (galectin 1) (LGALS1), mRNA /cds=(69,476) /gb=NM_002305 /gi=6006015 /ug=Hs.382367 /len=526	NM_002305	Hs.382367
4349	0.041391	cytokine receptor-like factor 1 (CRLF1), mRNA	NM_004750	Hs.114948
4395	0.047	hypothetical protein LOC51255 (LOC51255), mRNA /cds=(31,492) /gb=NM_016494 /gi=24475978 /ug=Hs.11156 /len=601	NM_016494	Hs.11156
4419	0.00215	BAC clone RP11-592A2 from 2, complete sequence	AC068288	
4420	5.88E-04	Machado-Joseph disease (spinocerebellar ataxia 3, olivopontocerebellar ataxia 3, autosomal dominant, ataxin 3) (MJD), transcript variant 1, mRNA /cds=(59,1144) /gb=NM_004993 /gi=13518018 /ug=Hs.66521 /len=1900	NM_004993	Hs.66521
4466	0.016439	cDNA: FLJ21659 fis, clone COL08743. /gb=AK025312 /gi=10437802 /ug=Hs.248862 /len=2423	AK025312	Hs.248862
4475	0.024176	RNA binding motif protein 8B (RBM8B) mRNA, complete cds	AF231512	
4542	0.014875	nudix (nucleoside diphosphate linked moiety X)-type motif 5 (NUDT5), mRNA /cds=(120,779) /gb=NM_014142 /gi=7657402 /ug=Hs.301957 /len=1129	NM_014142	Hs.301957



4549	0.01344	protein kinase, cAMP-dependent, regulatory, type I, alpha (tissue specific extinguisher 1) (PRKAR1A), mRNA /cds=(88,1233) /gb=NM_002734 /gi=4506062 /ug=Hs.183037 /len=3036	NM_002734	Hs.183037
4551	0.041391	mitochondrion, complete genome	NC_001807	
4554	0.01344	integral membrane protein 2B (ITM2B), mRNA /cds=(171,971) /gb=NM_021999 /gi=11527401 /ug=Hs.239625 /len=1843	NM_021999	Hs.239625
4571	0.017272	chromosome 1 open reading frame 13 (C1orf13), mRNA /cds=(45,1007) /gb=NM_030769 /gi=13540532 /ug=Hs.23756 /len=1552	NM_030769	Hs.23756
4602	0.009316	FLJ23277 protein (FLJ23277), mRNA /cds=(141,3089) /gb=NM_032236 /gi=18860906 /ug=Hs.334477 /len=3911	NM_032236	Hs.334477
4614	0.033276	repetitive sequence		
4694	0.005343	hypoxia-inducible factor 1, alpha subunit (basic helix-loop-helix transcription factor) (HIF1A), mRNA /cds=(265,2745) /gb=NM_001530 /gi=4504384 /ug=Hs.197540 /len=3933	NM_001530	Hs.197540
4709	0.037968	Alu repeat mRNA sequence /gb=AF330042 /gi=17225569 /ug=Hs.133319 /len=1440	AF330042	Hs.133319
4739	0.012127	ubiquitin carboxyl-terminal esterase L3 (ubiquitin thiolesterase) (UCHL3), mRNA /cds=(43,735) /gb=NM_006002 /gi=20149578 /ug=Hs.77917 /len=911	NM_006002	Hs.77917
4756	0.031823	chromobox 3 (HP1 gamma Drosophila) (CBX3), transcript variant 2, mRNA /cds=(152,703) /gb=NM_016587 /gi=20544150 /ug=Hs.406384 /len=1851	NM_016587	Hs.406384
4760	0.012769	BTB and CNC 1, basic leucine zipper transcription factor 1 (BACH1), mRNA /cds=(119,2329) /gb=NM_001186 /gi=4502352 /ug=Hs.154276 /len=5475	NM_001186	Hs.154276
4761	0.024176	calpastatin (CAST), transcript variant 2, mRNA /cds=(155,2215) /gb=NM_173060 /gi=27765084 /ug=Hs.359682 /len=4296	NM_173060	Hs.359682
4836	0.016439	nuclear receptor coactivator 3 (NCOA3), mRNA /cds=(184,4422) /gb=NM_006534 /gi=5729725 /ug=Hs.225977 /len=6754	NM_006534	Hs.225977
4884	0.02533	geranylgeranyl diphosphate synthase 1 (GGPS1), mRNA /cds=(233,1135) /gb=NM_004837 /gi=21359876 /ug=Hs.55498 /len=1489	NM_004837	Hs.55498
4885	0.006705	clone alpha_est218/52C1 mRNA sequence /gb=AF001542 /gi=2529714 /ug=Hs.356442 /len=2992	AF001542	Hs.356442
4915	0.01564	hypothetical protein FLJ13149 (FLJ13149), mRNA /cds=(291,2585) /gb=NM_021826 /gi=11141902 /ug=Hs.112188 /len=2836	NM_021826	Hs.112188
4950	0.006339	mRNA for KIAA1865 protein, partial cds. /cds=(622,2793) /gb=AB058768 /gi=14017946 /ug=Hs.179260 /len=3641	AB058768	Hs.179260
4976	0.01344	Sec23 A (S. cerevisiae), clone MGC:26267 IMAGE:4821858, mRNA, complete cds	BC036649	Hs.272927

4984	0.024176	LUC7-like ( <i>S. cerevisiae</i> ) (LUC7L), mRNA /cds=(89,1066) /gb=NM_018032 /gi=21359922 /ug=Hs.16803 /len=1542	NM_018032	Hs.16803
4990	0.041391	oxidation resistance 1 (OXR1), mRNA /cds=(31,840) /gb=NM_018002 /gi=8922240 /ug=Hs.169111 /len=1760	NM_018002	Hs.169111
5018	0.018141	mRNA; cDNA DKFZp762B195 (from clone DKFZp762B195) /gb=AL359585 /gi=8655645 /ug=Hs.356766 /len=2183	AL359585	Hs.356766
5038	9.02E-04	hypothetical protein FLJ22649 similar to signal peptidase SPC22/23 (FLJ22649), mRNA /cds=(114,656) /gb=NM_021928 /gi=11345461 /ug=Hs.42194 /len=1544	NM_021928	Hs.42194
5112	0.01564	inhibitor of DNA binding 1, dominant negative helix-loop-helix protein (ID1), mRNA /cds=(36,500) /gb=NM_002165 /gi=4504568 /ug=Hs.75424 /len=926	NM_002165	Hs.75424
5121	0.027779	hypothetical protein FLJ23436 (FLJ23436), mRNA /cds=(237,1799) /gb=NM_024671 /gi=20127628 /ug=Hs.85658 /len=2324	NM_024671	Hs.85658
5146	3.01E-05	creatine kinase, muscle (CKM), mRNA /cds=(75,1220) /gb=NM_001824 /gi=21536287 /ug=Hs.334347 /len=1620	NM_001824	Hs.334347
5154	0.003755	cullin 4A (CUL4A), mRNA /cds=(161,2140) /gb=NM_003589 /gi=11140810 /ug=Hs.183874 /len=3643	NM_003589	Hs.183874
5165	0.030424	additional sex combs like 1 ( <i>Drosophila</i> ) (ASXL1), mRNA /cds=(259,4884) /gb=NM_015338 /gi=27734730 /ug=Hs.3686 /len=6864	NM_015338	Hs.3686
5182	0.045065	C-type lectin BIMLEC precursor (BIMLEC), mRNA /cds=(12,710) /gb=NM_014880 /gi=26892292 /ug=Hs.2441 /len=1033	NM_014880	Hs.2441
5233	0.036347	ATP synthase, H transporting, mitochondrial F0 complex, subunit e (ATP5I), mRNA /cds=(64,273) /gb=NM_007100 /gi=6005716 /ug=Hs.85539 /len=336	NM_007100	Hs.85539
5234	0.031823	deiodinase, iodothyronine, type II (DIO2), transcript variant 1, mRNA /cds=(707,1528) /gb=NM_013989 /gi=7549802 /ug=Hs.154424 /len=6735	NM_013989	Hs.154424
5335	0.045065	hypothetical protein LOC284361 (LOC284361), mRNA /cds=(15,803) /gb=NM_175063 /gi=28372572 /ug=Hs.250465 /len=1907	NM_175063	Hs.250465
5349	0.008362	peroxisome receptor 1 (PXR1), mRNA /cds=(52,1947) /gb=NM_000319 /gi=21361203 /ug=Hs.158084 /len=3227	NM_000319	Hs.158084
5485	0.02533	mRNA for KIAA1896 protein, partial cds. /cds=(206,1912) /gb=AB067483 /gi=15620850 /ug=Hs.5476 /len=3712	AB067483	Hs.5476
5576	0.020975	mRNA; cDNA DKFZp451A0419 (from clone DKFZp451A0419) /gb=AL833070 /gi=21733661 /ug=Hs.146233 /len=4831	AL833070	Hs.146233

5578	0.045065	ATP synthase, H transporting, mitochondrial F1 complex, beta polypeptide (ATP5B), nuclear gene encoding mitochondrial protein, mRNA /cds=(46,1665) /gb=NM_001686 /gi=4502294 /ug=Hs.406510 /len=1807	NM_001686	Hs.406510
5585	0.031823	PHD finger protein 2 (PHF2), transcript variant 1, mRNA /cds=(27,3332) /gb=NM_005392 /gi=24797092 /ug=Hs.93868 /len=5279	NM_005392	Hs.93868
5609	0.045065	peroxisomal proliferator-activated receptor A interacting complex 285 (PRIC285), mRNA /cds=(425,6667) /gb=NM_033405 /gi=21703357 /ug=Hs.151714 /len=7804	NM_033405	Hs.151714
5611	0.009828	RNA binding motif protein 10 (RBM10), transcript variant 1, mRNA /cds=(380,3172) /gb=NM_005676 /gi=23111016 /ug=Hs.154583 /len=3415	NM_005676	Hs.154583
5677	0.005043	milk fat globule-EGF factor 8 protein (MFGE8), mRNA /cds=(61,1224) /gb=NM_005928 /gi=5174556 /ug=Hs.3745 /len=1934	NM_005928	Hs.3745
5719	0.00709	amplified in osteosarcoma (OS-9), mRNA /cds=(86,2089) /gb=NM_006812 /gi=5803108 /ug=Hs.76228 /len=2736	NM_006812	Hs.76228
5732	0.004758	cDNA FLJ37084 fis, clone BRACE2016583. /gb=AK094403 /gi=21753457 /ug=Hs.370346 /len=2151	AK094403	Hs.370346
5805	0.037968	hypothetical protein FLJ10583 (FLJ10583), mRNA /cds=(158,1387) /gb=NM_018148 /gi=8922537 /ug=Hs.105633 /len=2478	NM_018148	Hs.105633
5807	0.009828	mRNA for KIAA2019 protein. /cds=(15,8408) /gb=AB095939 /gi=24899201 /ug=Hs.57548 /len=9217	AB095939	Hs.57548
5811	0.029076	cDNA FLJ31038 fis, clone HSYRA2000159. /gb=AK055600 /gi=16550370 /ug=Hs.303154 /len=2981	AK055600	Hs.303154
5839	0.018141	ADP-ribosylation factor-like 5 (ARL5), mRNA /cds=(67,606) /gb=NM_012097 /gi=6912243 /ug=Hs.342849 /len=1281	NM_012097	Hs.342849
5843	0.045065	MYE4197a Myeloma (MYE) cDNA library cDNA, mRNA sequence /gb=BF174993 /gi=13441207 /ug=Hs.332023 /len=338	BF174993	Hs.332023
5858	0.026531	catenin (cadherin-associated protein), delta 1 (CTNND1), mRNA /cds=(537,3362) /gb=NM_001331 /gi=10835009 /ug=Hs.166011 /len=6232	NM_001331	Hs.166011
5869	0.007493	ATPase, Class I, type 8B, member 1 (ATP8B1), mRNA /cds=(1,3756) /gb=NM_005603 /gi=5031696 /ug=Hs.406187 /len=3756	NM_005603	Hs.406187
5876	0.037968	COX15 cytochrome c oxidase assembly protein (yeast) (COX15), nuclear gene encoding mitochondrial protein, transcript variant 2, mRNA /cds=(52,1218) /gb=NM_004376 /gi=17921986 /ug=Hs.226581 /len=2841	NM_004376	Hs.226581

5878	0.024176	zinc finger protein 36, C3H type, (mouse) (ZFP36), mRNA /cds=(60,1040) /gb=NM_003407 /gi=4507960 /ug=Hs.343586 /len=1746	NM_003407	Hs.343586
5908	0.029076	no match		
6017	0.036347	growth arrest-specific 7 (GAS7), transcript variant b, mRNA /cds=(122,1360) /gb=NM_005890 /gi=5360211 /ug=Hs.226133 /len=7979	NM_005890	Hs.226133
6054	0.043196	poly(A) binding protein, cytoplasmic 1 (PABPC1), mRNA /cds=(503,2404) /gb=NM_002568 /gi=4505574 /ug=Hs.172182 /len=2848	NM_002568	Hs.172182
6066	0.02533	actin related protein 2/3 complex, subunit 2, 34kDa (ARPC2), transcript variant 1, mRNA /cds=(113,1015) /gb=NM_152862 /gi=23238210 /ug=Hs.83583 /len=1462	NM_152862	Hs.83583
6073	0.029076	tuberous sclerosis 1 (TSC1), mRNA /cds=(222,3716) /gb=NM_000368 /gi=24475626 /ug=Hs.79393 /len=8600	NM_000368	Hs.79393
6077	0.026531	vaccinia related kinase 2 (VRK2), mRNA /cds=(131,1657) /gb=NM_006296 /gi=5454163 /ug=Hs.82771 /len=1833	NM_006296	Hs.82771
6113	0.037968	low density lipoprotein receptor (familial hypercholesterolemia) (LDLR), mRNA /cds=(94,2676) /gb=NM_000527 /gi=8051613 /ug=Hs.213289 /len=5175	NM_000527	Hs.213289
6121	0.041391	nm53h08.s1 NCI_CGAP_Br2 cDNA clone IMAGE:1071999 3' similar to gb:M10036 TRIOSEPHOSPHATE ISOMERASE mRNA sequence /clone=IMAGE:1071999 /clone_end=3' /gb=AA573489 /gi=2348017 /ug=Hs.404623 /len=230	AA573489	Hs.404623
6135	0.045065	cyclin-dependent kinase 4 (CDK4), transcript variant 1, mRNA /cds=(228,1139) /gb=NM_000075 /gi=16936531 /ug=Hs.95577 /len=1474	NM_000075	Hs.95577
6183	0.030424	actin related protein 2/3 complex, subunit 1B, 41kDa (ARPC1B), mRNA /cds=(90,1208) /gb=NM_005720 /gi=22907055 /ug=Hs.433506 /len=1520	NM_005720	Hs.433506
6247	0.001891	interleukin enhancer binding factor 3, 90kDa (ILF3), transcript variant 1, mRNA /cds=(267,2951) /gb=NM_012218 /gi=24234749 /ug=Hs.256583 /len=6058	NM_012218	Hs.256583
6252	0.039649	ribosomal protein S11 (RPS11), mRNA /cds=(34,510) /gb=NM_001015 /gi=14277698 /ug=Hs.182740 /len=594	NM_001015	Hs.182740
6254	0.041391	utrophin to dystrophin) (UTRN), mRNA /cds=(1,10302) /gb=NM_007124 /gi=6005937 /ug=Hs.286161 /len=10302	NM_007124	Hs.286161
6273	0.011512	clone IMAGE:4214654, mRNA	BC035518	Hs.16193
6290	0.033276	mRNA; cDNA DKFZp547C1510 (from clone DKFZp547C1510) /gb=AL831873 /gi=21732367 /ug=Hs.377056 /len=2405	AL831873	Hs.377056
6347	0.010364	mitochondrion, complete genome	NC_001807	

6348	0.005343	major histocompatibility complex, class I, F (HLA-F), mRNA /cds=(1,1089) /gb=NM_018950 /gi=9665231 /ug=Hs.110309 /len=1188	NM_018950	Hs.110309
6358	0.005498	ATP synthase, H transporting, mitochondrial FO complex, subunit f, isoform 2 (ATP5J2), mRNA /cds=(28,312) /gb=NM_004889 /gi=4757811 /ug=Hs.235557 /len=452	NM_004889	Hs.235557
6386	0.045065	mitochondrion, complete genome	NC_001807	
6391	0.018141	GABA(A) receptor-associated protein like 1 (GABARAPL1), mRNA /cds=(282,635) /gb=NM_031412 /gi=13899218 /ug=Hs.336429 /len=1933	NM_031412	Hs.336429
6403	0.003535	glyceronephosphate O-acyltransferase (GNPAT), mRNA /cds=(158,2200) /gb=NM_014236 /gi=7657133 /ug=Hs.12482 /len=2470	NM_014236	Hs.12482
6407	0.002292	hypothetical protein LOC51244 (LOC51244), mRNA /cds=(340,1233) /gb=NM_016474 /gi=24475969 /ug=Hs.158006 /len=1614	NM_016474	Hs.158006
6443	0.005658	Mlx interactor (MONDOA), mRNA /cds=(153,1733) /gb=NM_014938 /gi=7662347 /ug=Hs.52081 /len=4339	NM_014938	Hs.52081
6447	0.020973	vitamin A responsive; cytoskeleton related (JWA), mRNA /cds=(90,656) /gb=NM_006407 /gi=7669496 /ug=Hs.92384 /len=2088	NM_006407	Hs.92384
6468	0.024176	BAC clone RP11-629N4 from 4, complete sequence	AC079995	
6476	0.049002	cDNA: FLJ22425 fis, clone HRC08686	AK026078	Hs.288555
6515	0.016439	defensin, beta 1 (DEFB1), mRNA /cds=(72,278) /gb=NM_005218 /gi=13124884 /ug=Hs.32949 /len=366	NM_005218	Hs.32949
6530	0.017272	pleckstrin domain containing, family A (phosphoinositide binding specific) member 4 (PLEKHA4), mRNA /cds=(526,2865) /gb=NM_020904 /gi=10190743 /ug=Hs.9469 /len=3056	NM_020904	Hs.9469
6555	0.023066	U6 snRNA-associated Sm-like protein LSM8 (LSM8), mRNA /cds=(63,353) /gb=NM_016200 /gi=21314665 /ug=Hs.241578 /len=1206	NM_016200	Hs.241578
6558	0.006705	ribosomal protein S23 (RPS23), mRNA /cds=(32,463) /gb=NM_001025 /gi=14790142 /ug=Hs.3463 /len=506	NM_001025	Hs.3463
6569	0.041391	tuberous sclerosis 2 (TSC2), transcript variant 1, mRNA /cds=(19,5442) /gb=NM_000548 /gi=10938006 /ug=Hs.90303 /len=5543	NM_000548	Hs.90303
6583	0.029076	KIAA0170 gene product (KIAA0170), mRNA /cds=(14,6283) /gb=NM_014641 /gi=7661965 /ug=Hs.433653 /len=6940	NM_014641	Hs.433653
6596	0.004758	dual specificity phosphatase 5 (DUSP5), mRNA /cds=(211,1365) /gb=NM_004419 /gi=12707565 /ug=Hs.2128 /len=2473	NM_004419	Hs.2128
6656	0.029076	protein phosphatase 1, regulatory (inhibitor) subunit 2 (PPP1R2), mRNA /cds=(235,852) /gb=NM_006241 /gi=19923357 /ug=Hs.267819 /len=3355	NM_006241	Hs.267819
6663	0.02533	PRO0461 protein (PRO0461), mRNA /gb=NM_031268 /gi=20588827 /ug=Hs.25063 /len=1100	NM_031268	Hs.25063

6678	0.009828	vesicle-associated membrane protein 8 (endobrevin) (VAMP8), mRNA /cds=(54,356) /gb=NM_003761 /gi=14043025 /ug=Hs.172684 /len=702	NM_003761	Hs.172684
6720	0.01344	nascent-polypeptide-associated complex alpha polypeptide (NACA), mRNA /cds=(26,673) /gb=NM_005594 /gi=5031930 /ug=Hs.32916 /len=797	NM_005594	Hs.32916
6775	0.039649	protein tyrosine phosphatase type IVA, member 2 (PTP4A2), transcript variant 1, mRNA /cds=(1011,1514) /gb=NM_003479 /gi=18104974 /ug=Hs.82911 /len=3925	NM_003479	Hs.82911
6796	0.021999	hypothetical protein, clone 2746033 (HSA272196), mRNA /cds=(39,593) /gb=NM_018405 /gi=24475639 /ug=Hs.8179 /len=861	NM_018405	Hs.8179
6814	0.027779	regulator of G-protein signalling 5 (RGS5), mRNA /cds=(82,627) /gb=NM_003617 /gi=4506518 /ug=Hs.24950 /len=2076	NM_003617	Hs.24950
6843	0.008828	mannosidase, alpha, class 1A, member 1 (MAN1A1), mRNA /cds=(443,2404) /gb=NM_005907 /gi=24497518 /ug=Hs.25253 /len=4139	NM_005907	Hs.25253
6844	0.031823	hypothetical protein CLONE24945 (CLONE24945), mRNA /cds=(144,1367) /gb=NM_015683 /gi=18373304 /ug=Hs.30882 /len=2518	NM_015683	Hs.30882
6886	0.00313	inhibitor of Bruton's tyrosine kinase (IBTK), mRNA /cds=(420,1031) /gb=NM_015525 /gi=24308082 /ug=Hs.306425 /len=2240	NM_015525	Hs.306425
6892	0.005658	hypothetical protein FLJ10420 (FLJ10420), mRNA /cds=(34,825) /gb=NM_018090 /gi=20127581 /ug=Hs.289087 /len=2046	NM_018090	Hs.289087
6898	0.002017	plexin B2 (PLXNB2), mRNA /cds=(26,1438) /gb=NM_012401 /gi=20270189 /ug=Hs.3989 /len=2175	NM_012401	Hs.3989
6900	0.037968	developmentally regulated GTP binding protein 1 (DRG1), mRNA /cds=(66,1169) /gb=NM_004147 /gi=4758795 /ug=Hs.115242 /len=1383	NM_004147	Hs.115242
6922	0.020975	cartilage acidic protein 1 (CRTAC1), mRNA /cds=(319,1575) /gb=NM_018058 /gi=8922351 /ug=Hs.326444 /len=2178	NM_018058	Hs.326444
7055	0.011512	CDC7 cell division cycle 7-like 1 (S. cerevisiae) (CDC7L1), mRNA /cds=(126,1850) /gb=NM_003503 /gi=11038647 /ug=Hs.28853 /len=3178	NM_003503	Hs.28853
7074	0.039649	microsomal glutathione S-transferase 3 (MGST3), mRNA /cds=(89,547) /gb=NM_004528 /gi=22035640 /ug=Hs.424468 /len=688	NM_004528	Hs.424468
7078	0.031823	KIAA0438 gene product (KIAA0438), mRNA /cds=(118,2244) /gb=NM_014819 /gi=7662123 /ug=Hs.279849 /len=4765	NM_014819	Hs.279849
7103	0.045065	sialidase 1 (lysosomal sialidase) (NEU1), mRNA /cds=(130,1377) /gb=NM_000434 /gi=4557790 /ug=Hs.118721 /len=1894	NM_000434	Hs.118721
7169	0.041391	GDP dissociation inhibitor 2 (GDI2), mRNA /cds=(153,1490) /gb=NM_001494 /gi=6598322 /ug=Hs.56845 /len=2274	NM_001494	Hs.56845

7181	0.018141	eukaryotic translation initiation factor 4A, isoform 1 (EIF4A1), mRNA /cds=(17,1237) /gb=NM_001416 /gi=4503528 /ug=Hs.129673 /len=1383	NM_001416	Hs.129673
7193	0.007917	heterogeneous nuclear ribonucleoprotein A1 (HNRPA1), transcript variant 2, mRNA /cds=(105,1223) /gb=NM_031157 /gi=14043069 /ug=Hs.376844 /len=1925	NM_031157	Hs.376844
7196	0.01564	ubiquitin B (UBB), mRNA /cds=(139,828) /gb=NM_018955 /gi=22538474 /ug=Hs.356190 /len=971	NM_018955	Hs.356190
7223	0.019047	myosin, light polypeptide 6, alkali, smooth muscle and non-muscle (MYL6), transcript variant 3, mRNA /cds=(41,514) /gb=NM_079425 /gi=17986263 /ug=Hs.77385 /len=717	NM_079425	Hs.77385
7262	0.033276	LIM and senescent cell antigen-like domains 1 (LIMS1), mRNA /cds=(120,1097) /gb=NM_004987 /gi=13518025 /ug=Hs.112378 /len=1236	NM_004987	Hs.112378
7284	0.01344	GK003 protein (GK003), mRNA /cds=(10,690) /gb=NM_020192 /gi=21281666 /ug=Hs.83313 /len=901	NM_020192	Hs.83313
7303	0.018141	cDNA FLJ38955 fis, clone NT2RI2000107. /gb=AK096274 /gi=21755725 /ug=Hs.156100 /len=2613	AK096274	Hs.156100
7371	0.027779	hypothetical protein MGC14156 (MGC14156), mRNA /cds=(83,427) /gb=NM_032906 /gi=14249679 /ug=Hs.26136 /len=1325	NM_032906	Hs.26136
7374	0.031823	sarcoglycan, epsilon (SGCE), mRNA /cds=(69,1382) /gb=NM_003919 /gi=10835046 /ug=Hs.110708 /len=1658	NM_003919	Hs.110708
7406	0.00709	hypothetical protein FLJ14511 (FLJ14511), mRNA /cds=(23,1273) /gb=NM_033087 /gi=14861835 /ug=Hs.40919 /len=1835	NM_033087	Hs.40919
7509	0.043196	cDNA FLJ12900 fis, clone NT2RP2004321. /gb=AK022962 /gi=10434656 /ug=Hs.21851 /len=2603	AK022962	Hs.21851
7568	0.014142	te45b07.x1 Soares_NhHMPu_S1 cDNA clone IMAGE:2089621 3', mRNA sequence /clone=IMAGE:2089621 /clone_end=3' /gb=AI554849 /gi=4487212 /ug=Hs.171343 /len=239	AI554849	Hs.171343
7578	0.009828	hypothetical protein DKFZp586G0123 (DKFZp586G0123), mRNA /cds=(25,315) /gb=NM_013386 /gi=9558726 /ug=Hs.24713 /len=1294	NM_013386	Hs.24713
7614	0.008828	ALS2CR3 gene for amyotrophic lateral sclerosis 2, candidate 3, exon 14	AB038962	
7622	0.012769	mRNA for KIAA1912 protein, partial cds. /cds=(395,2164) /gb=AB067499 /gi=15620882 /ug=Hs.283902 /len=3985	AB067499	Hs.283902
7655	0.020975	dihydropyrimidinase-like 2 (DPYSL2), mRNA /cds=(275,1993) /gb=NM_001386 /gi=19923654 /ug=Hs.173381 /len=4459	NM_001386	Hs.173381

7677	0.009828	uveal autoantigen with coiled-coil domains and ankyrin repeats (UACA), mRNA /cds=(50,4300) /gb=NM_018003 /gi=24308154 /ug=Hs.49753 /len=4428	NM_018003	Hs.49753
7681	0.024176	likely ortholog of mouse IRA1 protein (IRA1), mRNA /cds=(227,1771) /gb=NM_024665 /gi=19913370 /ug=Hs.350547 /len=3950	NM_024665	Hs.350547
7712	0.036347	Janus kinase 2 (a protein tyrosine kinase) (JAK2), mRNA /cds=(495,3893) /gb=NM_004972 /gi=13325062 /ug=Hs.115541 /len=5097	NM_004972	Hs.115541
7718	0.047	echinoderm microtubule associated protein like 1 (EML1), mRNA /cds=(363,2516) /gb=NM_004434 /gi=4758267 /ug=Hs.12451 /len=3962	NM_004434	Hs.12451
7835	0.003327	BAC clone RP11-182E7 from 2, complete sequence	AC073255	
7856	0.043196	hypothetical protein MGC12103 (LOC133619), mRNA /cds=(142,1479) /gb=NM_130809 /gi=18677734 /ug=Hs.6614 /len=1895	NM_130809	Hs.6614
7863	0.034783	mRNA; cDNA DKFZp686A1444 (from clone DKFZp686A1444)	AL833555	Hs.278428
7870	0.021999	BX107856 Soares fetal liver spleen 1NFLS cDNA clone IMAGp998A13537, mRNA sequence /clone=IMAGp998A13537_/_IMAGE:248340 /gb=BX107856 /gi=27834894 /ug=Hs.329327 /len=736	BX107856	Hs.329327
7902	0.045065	cDNA FLJ40622 fis, clone THYMU2013779. /gb=AK097941 /gi=21757847 /ug=Hs.374352 /len=1754	AK097941	Hs.374352
7936	0.024176	UI-H-BI4-apg-d-10-0-UI.s1 NCI_CGAP_Sub8 cDNA clone IMAGE:3087402 3', mRNA sequence /clone=IMAGE:3087402 /clone_end=3' /gb=BF509764 /gi=11593062 /ug=Hs.439798 /len=1099	BF509764	Hs.439798
7940	0.045065	UI-CF-DU1-adl-j-09-0-UI.s1 UI-CF-DU1 cDNA clone UI-CF-DU1-adl-j-09-0-UI 3', mRNA sequence /clone=UI-CF-DU1-adl-j-09-0-UI /clone_end=3' /gb=BM979130 /gi=19599265 /ug=Hs.114675 /len=745	BM979130	Hs.114675
7943	0.019047	DNA sequence from clone RP11-490F3 on chromosome 9 Contains a cytochrome C pseudogene, 2 isoforms of the 5' end of a gene for a novel protein and a CpG island, complete sequence	AL360020	
7944	0.039649	smad-interacting protein-1 gene, partial cds	AY029472	
7984	0.007917	hypothetical protein FLJ11126 (FLJ11126), mRNA /cds=(81,1517) /gb=NM_018332 /gi=20070297 /ug=Hs.226396 /len=2927	NM_018332	Hs.226396
7993	0.029076	antigen identified by monoclonal antibody Ki-67 (MKI67), mRNA /cds=(197,9967) /gb=NM_002417 /gi=19923216 /ug=Hs.80976 /len=12515	NM_002417	Hs.80976
8047	0.010364	oxysterol binding protein-like 9 (OSBPL9), transcript variant 7, mRNA /cds=(20,2260) /gb=NM_148909 /gi=22547175 /ug=Hs.21938 /len=2949	NM_148909	Hs.21938
8096	0.036347	thymosin, beta 10 (TMSB10), mRNA /cds=(66,200) /gb=NM_021103 /gi=10863894 /ug=Hs.76293 /len=453	NM_021103	Hs.76293



8114	0.00111	URB mRNA, complete cds /cds=(146,2998) /gb=AF506819 /gi=21039408 /ug=Hs.356289 /len=3320	AF506819	Hs.356289
8175	0.010364	cDNA FLJ40228 fis, clone TESTI2022623, moderately similar to Rattus norvegicus KPL2 (Kpl2) mRNA	AK097547	
8182	0.036347	general transcription factor IIH, polypeptide 2, 44kDa (GTF2H2), mRNA /cds=(1,1188) /gb=NM_001515 /gi=6681761 /ug=Hs.191356 /len=1188	NM_001515	Hs.191356
8206	0.045065	mRNA for KIAA0640 protein, partial cds. /cds=(1,1813) /gb=AB014540 /gi=3327093 /ug=Hs.153026 /len=4824	AB014540	Hs.153026
8227	0.031823	hypothetical protein FLJ20628 (FLJ20628), mRNA /cds=(23,1456) /gb=NM_017910 /gi=13435382 /ug=Hs.32356 /len=1846	NM_017910	Hs.32356
8298	0.006339	chromosome 14 DNA sequence BAC R-159L20 of library RPCI-11 from chromosome 14 of complete sequence	AL121852	
8306	0.004758	hypothetical protein FLJ20313 (FLJ20313), mRNA /cds=(345,1700) /gb=NM_017762 /gi=8923296 /ug=Hs.126721 /len=2226	NM_017762	Hs.126721
8328	0.043196	asporin (LRR class 1) (ASPN), mRNA	NM_017680	Hs.10760
8336	0.024176	cDNA: FLJ20913 fis, clone ADSE00630. /gb=AK024566 /gi=10436876 /ug=Hs.7063 /len=1268	AK024566	Hs.7063
8360	0.036347	chromosome 14 DNA sequence BAC R-103F13 of library RPCI-11 from chromosome 14 of complete sequence	AL359240	
8372	0.009828	mRNA; cDNA DKFZp547M059 (from clone DKFZp547M059) /gb=AL831946 /gi=21732473 /ug=Hs.433066 /len=2922	AL831946	Hs.433066
8382	0.039649	hypothetical protein MGC5149 (MGC5149), mRNA	XM_051200	
8402	0.041391	BAC clone RP11-326O23 from 4, complete sequence	AC096740	
8404	0.034783	cDNA FLJ11049 fis, clone PLACE1004548	AK001911	Hs.107287
8418	0.020975	chromosome 5 clone CTB-162J7, complete sequence	AC010297	
8479	9.02E-04	clone IMAGE:5259731, mRNA /gb=BC033052 /gi=22766860 /ug=Hs.284707 /len=1340	BC033052	Hs.284707
8490	0.031823	chromosome 11, clone CTD-2010I16, complete sequence	AC084337	
8494	0.018141	UI-H-DH0-aui-j-10-0-UI.s1 NCI_CGAP_DH0 cDNA clone IMAGE:5871081 3', mRNA sequence /clone=IMAGE:5871081 /clone_end=3' /gb=BM994461 /gi=19719362 /ug=Hs.434057 /len=2059	BM994461	Hs.434057
8509	0.034783	pericentrin 1 (PCNT1), mRNA /cds=(81,2051) /gb=NM_024844 /gi=13376258 /ug=Hs.184352 /len=2147	NM_024844	Hs.184352
8553	0.007917	DNA sequence from clone RP4-765C7 on chromosome 1, complete sequence	AL365357	
8555	0.017272	genomic DNA, chromosome 11q clone:RP11-762B21, complete sequence	AP000926	
8568	0.023066	clone IMAGE:4825278, mRNA	BC034789	Hs.323849

8623	0.02533	cDNA FLJ31753 fis, clone NT2RI2007468. /gb=AK056315 /gi=16551681 /ug=Hs.349283 /len=2361	AK056315	Hs.349283
8624	0.019047	mRNA; cDNA DKFZp313C0935 (from clone DKFZp313C0935) /gb=AL832706 /gi=21733285 /ug=Hs.433110 /len=3270	AL832706	Hs.433110
8659	0.026531	BAC clone RP11-363G15 from 4, complete sequence	AC096746	
8662	0.023066	ts93d11.x1 NCI_CGAP_GC6 cDNA clone IMAGE:2238837 3', mRNA sequence /clone=IMAGE:2238837 /clone_end=3' /gb=AI631165 /gi=4682495 /ug=Hs.196952 /len=537	AI631165	Hs.196952
8677	0.003755	DNA sequence from clone RP11-185C19 on chromosome 1q31.1-31.3, complete sequence	AL139133	
8684	0.031823	chromosome 3 clone RP11-321A23, complete sequence	AC096970	
8686	0.00709	mRNA; cDNA DKFZp564P016 (from clone DKFZp564P016) /gb=AL049337 /gi=4500118 /ug=Hs.132571 /len=1938	AL049337	Hs.132571
8687	0.043196	chromosome 5 clone CTD-2337D22, complete sequence	AC008952	
8709	0.039649	clone alpha_est218/52C1 mRNA sequence /gb=AF001542 /gi=2529714 /ug=Hs.356442 /len=2992	AF001542	Hs.356442
8710	0.033276	alpha gene sequence	AF203815	
8722	0.043196	EST388508 MAGE resequences, MAGN cDNA, mRNA sequence /gb=AW976399 /gi=8167626 /ug=Hs.293205 /len=718	AW976399	Hs.293205
8744	6.33E-04	we12e11.x1 NCI_CGAP_Lu24 cDNA clone IMAGE:2340908 3', mRNA sequence /clone=IMAGE:2340908 /clone_end=3' /gb=AI912685 /gi=5632540 /ug=Hs.213734 /len=567	AI912685	Hs.213734
8745	0.023066	chromosome 18, clone RP11-49I11, complete sequence	AC023043	
8779	0.018141	thioredoxin-like 2 (TXNL2), mRNA /cds=(5,1012) /gb=NM_006541 /gi=5730103 /ug=Hs.42644 /len=1942	NM_006541	Hs.42644
8780	0.009828	MR2-CI0186-291100-010-a06 CI0186 cDNA, mRNA sequence /gb=BF814502 /gi=12147047 /ug=Hs.446594 /len=530	BF814502	Hs.446594
8995	0.030424	EST1000 HEV PCR-select cDNA clone HEV#2154, mRNA sequence /clone=HEV#2154 /gb=BM956063 /gi=23346294 /ug=Hs.390155 /len=664	BM956063	Hs.390155
9028	0.037968	UI-H-CO0-arf-f-05-0-UI.s1 NCI_CGAP_Sub9 cDNA clone IMAGE:3106304 3', mRNA sequence /clone=IMAGE:3106304 /clone_end=3' /gb=BQ027925 /gi=19763204 /ug=Hs.162459 /len=968	BQ027925	Hs.162459
9048	0.049002	SR rich protein (DKFZp564B0769), mRNA /cds=(33,2450) /gb=NM_032870 /gi=18699723 /ug=Hs.18368 /len=2663	NM_032870	Hs.18368
9063	0.009828	clone 23612 mRNA sequence /gb=U90902 /gi=1913880 /ug=Hs.82141 /len=1548	U90902	Hs.82141
9129	0.002599	alpha gene sequence	AF203815	

9132	0.017272	hypothetical protein LOC90410 (LOC90410), mRNA /cds=(142,588) /gb=NM_174887 /gi=28372522 /ug=Hs.4187 /len=1025	NM_174887	Hs.4187
9136	0.019991	UI-CF-EC1-aeb-n-22-0-UI.s1 UI-CF-EC1 cDNA clone UI-CF-EC1-aeb-n-22-0-UI 3', mRNA sequence /clone=UI-CF-EC1-aeb-n-22-0-UI /clone_end=3' /gb=BM978368 /gi=19597726 /ug=Hs.143789 /len=749	BM978368	Hs.143789
9143	0.01564	mRNA; cDNA DKFZp434N185 (from clone DKFZp434N185) /gb=AL117645 /gi=5912235 /ug=Hs.33032 /len=1641	AL117645	Hs.33032
9153	0.018141	AGENCOURT_8584280 Lupski_sympathetic_trunk cDNA clone IMAGE:6192820 5', mRNA sequence /clone=IMAGE:6192820 /clone_end=5' /gb=BQ876563 /gi=22268571 /ug=Hs.346743 /len=925	BQ876563	Hs.346743
9154	0.018141	mRNA; cDNA DKFZp564B213 (from clone DKFZp564B213) /gb=AL049240 /gi=4499973 /ug=Hs.380268 /len=767	AL049240	Hs.380268
9162	0.049002	ah98a04.s1 Soares_NFL_T_GBC_S1 cDNA clone IMAGE:1327086 3', mRNA sequence /clone=IMAGE:1327086 /clone_end=3' /gb=AA757363 /gi=2805226 /ug=Hs.105224 /len=444	AA757363	Hs.105224
9187	0.009828	LL5 beta (LL5beta), mRNA /cds=(116,3748) /gb=NM_145753 /gi=21955171 /ug=Hs.7378 /len=5491	NM_145753	Hs.7378
9189	0.045065	clone IMAGE:5265581, mRNA /gb=BC035165 /gi=23272508 /ug=Hs.400548 /len=2237	BC035165	Hs.400548
9201	0.041391	BAC clone RP11-798L22 from 2, complete sequence	AC093391	
9231	4.06E-04	tj44d11.x1 Soares_NSF_F8_9W_OT_PA_P_S1 cDNA clone IMAGE:2144373 3' similar to gb:Y00716 COMPLEMENT FACTOR H PRECURSOR mRNA sequence /clone=IMAGE:2144373 /clone_end=3' /gb=AI470482 /gi=4332572 /ug=Hs.387691 /len=384	AI470482	Hs.387691
9257	0.047	602246637F1 NIH_MGC_62 cDNA clone IMAGE:4331985 5', mRNA sequence /clone=IMAGE:4331985 /clone_end=5' /gb=BF690692 /gi=11976100 /ug=Hs.442332 /len=929	BF690692	Hs.442332
9270	0.003755	602122561F1 NIH_MGC_56 cDNA clone IMAGE:4279766 5', mRNA sequence /clone=IMAGE:4279766 /clone_end=5' /gb=BF668349 /gi=11942244 /ug=Hs.44731 /len=906	BF668349	Hs.44731
9325	0.023066	BAC clone RP11-730E1 from 4, complete sequence	AC093900	
9326	0.030424	UI-1-BB1p-akc-h-10-0-UI.s1 NCI_CGAP_PI6 cDNA clone UI-1-BB1p-akc-h-10-0-UI 3', mRNA sequence /clone=UI-1-BB1p-akc-h-10-0-UI /clone_end=3' /gb=BQ026175 /gi=19761454 /ug=Hs.127786 /len=1398	BQ026175	Hs.127786
9327	0.033276	mRNA; cDNA DKFZp566M063 (from clone DKFZp566M063)	AL110194	Hs.6727
9331	0.019047	cDNA FLJ31439 fis, clone NT2NE2000707. /gb=AK056001 /gi=16550873 /ug=Hs.349656 /len=2009	AK056001	Hs.349656

9400	0.031823	SKB1 (S. pombe) (SKB1), mRNA /cds=(92,2005) /gb=NM_006109 /gi=20070219 /ug=Hs.12912 /len=2413	NM_006109	Hs.12912
9408	0.037968	Down syndrome critical region gene 2 (DSCR2), mRNA /cds=(64,930) /gb=NM_003720 /gi=4505022 /ug=Hs.5198 /len=1080	NM_003720	Hs.5198
9482	0.023066	mRNA for KIAA1367 protein, partial cds. /cds=(1,1741) /gb=AB037788 /gi=7243114 /ug=Hs.224961 /len=4196	AB037788	Hs.224961
9543	0.033276	NADH dehydrogenase (ubiquinone) Fe-S protein 3, 30kDa (NADH-coenzyme Q reductase) (NDUFS3), mRNA /cds=(13,807) /gb=NM_004551 /gi=4758787 /ug=Hs.429506 /len=899	NM_004551	Hs.429506
9556	0.049002	hypothetical protein FLJ39155 (FLJ39155), mRNA /cds=(325,1536) /gb=NM_152403 /gi=22748856 /ug=Hs.20103 /len=2824	NM_152403	Hs.20103
9558	0.005658	Gene 33/Mig-6 (MIG-6), mRNA /cds=(213,1601) /gb=NM_018948 /gi=21314673 /ug=Hs.11169 /len=3099	NM_018948	Hs.11169
9564	0.030424	REV1-like (yeast) (REV1L), mRNA /cds=(213,3968) /gb=NM_016316 /gi=7706680 /ug=Hs.110347 /len=4276	NM_016316	Hs.110347
9626	0.026531	ATP-binding cassette, sub-family F (GCN20), member 1 (ABCF1), mRNA /cds=(95,2518) /gb=NM_001090 /gi=10947134 /ug=Hs.9573 /len=3141	NM_001090	Hs.9573
9662	0.019991	hypothetical protein FLJ33282 (FLJ33282), mRNA /cds=(225,1523) /gb=NM_152388 /gi=22748830 /ug=Hs.346509 /len=2078	NM_152388	Hs.346509
9745	0.036347	hypothetical protein MGC32104 (MGC32104), mRNA /cds=(101,1651) /gb=NM_144684 /gi=21389584 /ug=Hs.147025 /len=4732	NM_144684	Hs.147025
9806	0.01344	7k03e02.x1 NCI_CGAP_GC6 cDNA clone IMAGE:3443402 3', mRNA sequence /clone=IMAGE:3443402 /clone_end=3' /gb=BF056273 /gi=10810169 /ug=Hs.188920 /len=572	BF056273	Hs.188920
9836	0.019991	clone IMAGE:3629966, mRNA /gb=BC005082 /gi=13937698 /ug=Hs.334575 /len=1734	BC005082	Hs.334575
9840	0.043196	hypothetical protein FLJ20244 (FLJ20244), mRNA /cds=(89,2068) /gb=NM_017722 /gi=8923218 /ug=Hs.158947 /len=2137	NM_017722	Hs.158947
9850	0.020975	cDNA FLJ11946 fis, clone HEMBB1000709. /gb=AK022008 /gi=10433321 /ug=Hs.323231 /len=3241	AK022008	Hs.323231
9930	0.031823	cofactor required for Sp1 transcriptional activation, subunit 2, 150kDa (CRSP2), mRNA /cds=(120,4484) /gb=NM_004229 /gi=4758101 /ug=Hs.407604 /len=7984	NM_004229	Hs.407604
9944	0.047	CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5, EJ16, EJ30, EL32 and G344) (CD59), mRNA /cds=(50,436) /gb=NM_000611 /gi=20127410 /ug=Hs.278573 /len=1946	NM_000611	Hs.278573
9953	0.029076	fatty acid binding protein 1, liver (FABP1), mRNA	NM_001443	Hs.380135

10026	0.002017	dishevelled associated activator of morphogenesis 1 (DAAM1), mRNA /cds=(126,3362) /gb=NM_014992 /gi=21071076 /ug=Hs.197751 /len=4256	NM_014992	Hs.197751
10091	0.011507	zinc finger protein 258, clone IMAGE:5263804, mRNA	BC033903	Hs.410016
10193	0.045065	cDNA FLJ35309 fis, clone PROST2010219, weakly similar to CALMODULIN. /gb=AK092628 /gi=21751263 /ug=Hs.250861 /len=2364	AK092628	Hs.250861
10198	0.036347	ze65h12.s1 Soares retina N2b4HR cDNA clone IMAGE:363911 3', mRNA sequence /clone=IMAGE:363911 /clone_end=3' /gb=AA021186 /gi=1484920 /ug=Hs.226306 /len=422	AA021186	Hs.226306
10232	0.047	BX094256 Soares_fetal_heart_NbHH19W cDNA clone IMAGp998B20783, mRNA sequence /clone=IMAGp998B20783_ IMAGE:342835 /gb=BX094256 /gi=27841884 /ug=Hs.407356 /len=477	BX094256	Hs.407356
10245	0.031112	cDNA FLJ31147 fis, clone IMR322001438. /gb=AK055709 /gi=16550504 /ug=Hs.6670 /len=1747	AK055709	Hs.6670
10266	0.005043	hypothetical protein LOC282997 (LOC282997), mRNA	XM_208473	
10296	0.030424	cDNA FLJ11603 fis, clone HEMBA1003926	AK021665	
10340	0.039649	UI-E-CR1-aeb-c-08-0-UI.s1 UI-E-CR1 cDNA clone UI-E-CR1-aeb-c-08-0-UI 3', mRNA sequence /clone=UI-E-CR1-aeb-c-08-0-UI /clone_end=3' /gb=BU728348 /gi=23650137 /ug=Hs.437120 /len=1093	BU728348	Hs.437120
10359	0.037968	C1q and tumor necrosis factor related protein 2 (C1QTNF2), mRNA /cds=(130,987) /gb=NM_031908 /gi=19923636 /ug=Hs.110062 /len=2404	NM_031908	Hs.110062
10408	0.02533	yr31a03.r1 Soares fetal liver spleen 1NFLS cDNA clone IMAGE:206860 5' similar to contains MER19 repetitive element ;, mRNA sequence /clone=IMAGE:206860 /clone_end=5' /gb=R98895 /gi=985496 /ug=Hs.125823 /len=377	R98895	Hs.125823
10445	0.024176	mRNA for KIAA1715 protein, partial cds. /cds=(152,1441) /gb=AB051502 /gi=12697974 /ug=Hs.209561 /len=5714	AB051502	Hs.209561
10454	0.004758	genomic DNA, chromosome 11 clone:RP11-831H9, complete sequence	AP001458	
10465	0.020975	chromosome 19 clone CTC-425O23, complete sequence	AC118344	
10502	0.014142	clone IMAGE:5301489, mRNA	BC045690	Hs.6799
10523	0.024176	clone FLB6914 PRO1821 mRNA, complete cds (=AL050083.1)	AF130061	Hs.44143
10529	0.033276	df27e02.w1 Morton Fetal Cochlea cDNA clone IMAGE:2484578 3', mRNA sequence /clone=IMAGE:2484578 /clone_end=3' /gb=BI492664 /gi=15332008 /ug=Hs.345490 /len=657	BI492664	Hs.345490
10530	0.049002	EST383317 MAGE resequences, MAGL cDNA, mRNA sequence /gb=AW971229 /gi=8161074 /ug=Hs.293372 /len=642	AW971229	Hs.293372
10540	0.047	BX116063 NCI_CGAP_Brn23 cDNA clone IMAGp998O244959, mRNA sequence /clone=IMAGp998O244959_ IMAGE:2016239 /gb=BX116063 /gi=27839769 /ug=Hs.127872 /len=537	BX116063	Hs.127872

10581	0.023066	chromosome 5 clone CTB-113P19, complete sequence	AC011374	
10617	0.026531	SRY (sex determining region Y)-box 5 (SOX5), transcript variant B, mRNA /cds=(373,2625) /gb=NM_152989 /gi=23308714 /ug=Hs.87224 /len=4492	NM_152989	Hs.87224
10674	0.041385	EST54387 Fetal heart II cDNA 3' end, mRNA sequence /clone_end=3' /gb=AA347998 /gi=2000234 /ug=Hs.104574 /len=307	AA347998	Hs.104574
10693	0.027779	7f77g07.x1 Lupski_dorsal_root_ganglion cDNA clone IMAGE:3302989 3' similar to Q16465 HYPOTHETICAL PROTEIN ;, mRNA sequence /clone=IMAGE:3302989 /clone_end=3' /gb=BG057970 /gi=12523993 /ug=Hs.405856 /len=363	BG057970	Hs.405856
10703	0.047	chromosome 1 clone RP11-88N11, complete sequence	AC094022	
10724	0.047	no match		
10765	0.016439	BAC clone RP11-731119 from 2, complete sequence	AC093690	
10900	0.043196	UI-1-BC1p-asx-h-02-0-UI.s1 NCI_CGAP_PI3 cDNA clone UI-1-BC1p-asx-h-02-0-UI 3', mRNA sequence /clone=UI-1-BC1p-asx-h-02-0-UI /clone_end=3' /gb=BQ012708 /gi=19737609 /ug=Hs.191900 /len=590	BQ012708	Hs.191900
10918	0.010925	collagen, type V, alpha 2 (COL5A2), mRNA /cds=(158,4648) /gb=NM_000393 /gi=16554580 /ug=Hs.82985 /len=6217	NM_000393	Hs.82985
10930	0.026531	Similar to RIKEN cDNA 2810004N23 gene, clone MGC:46269 IMAGE:5589128, mRNA, complete cds /cds=(57,905) /gb=BC036800 /gi=22477333 /ug=Hs.390881 /len=1468	BC036800	Hs.390881
10956	0.011512	pp9974 mRNA, complete cds /cds=(2009,2350) /gb=AF318382 /gi=18027855 /ug=Hs.251664 /len=2630	AF318382	Hs.251664
10959	0.027779	7p65g03.x1 NCI_CGAP_Pr28 cDNA clone IMAGE:3650861 3', mRNA sequence /clone=IMAGE:3650861 /clone_end=3' /gb=BF436898 /gi=11449213 /ug=Hs.213352 /len=426	BF436898	Hs.213352
10976	0.033276	in56e04.x1 HR85 islet cDNA clone IMAGE:6126055 3', mRNA sequence /clone=IMAGE:6126055 /clone_end=3' /gb=BU784825 /gi=23830229 /ug=Hs.442971 /len=548	BU784825	Hs.442971
10999	0.024176	ax37a08.x1 Proliferating Erythroid Cells (LCB:ax library) cDNA clone ax37a08 random, mRNA sequence /clone=ax37a08 /gb=BG943384 /gi=14342756 /ug=Hs.339555 /len=555	BG943384	Hs.339555
11002	0.004487	AU119153 HEMBA1 cDNA clone HEMBA1005152 5', mRNA sequence /clone=HEMBA1005152 /clone_end=5' /gb=AU119153 /gi=10934388 /ug=Hs.288615 /len=820	AU119153	Hs.288615
11023	0.029076	cDNA FLJ36508 fis, clone TRACH2001121. /gb=AK093827 /gi=21752767 /ug=Hs.298250 /len=2182	AK093827	Hs.298250

11024	0.01344	hypothetical protein MGC33607 (MGC33607), mRNA /cds=(42,2543) /gb=NM_152775 /gi=22749514 /ug=Hs.41101 /len=2866	NM_152775	Hs.41101
11028	0.039649	similar to GA binding protein alpha chain (GABP-alpha subunit) (Transcription factor E4TF1-60) (Nuclear respiratory factor-2 subunit alpha) (LOC285899), mRNA	XM_209807	
11034	0.043196	mRNA; cDNA DKFZp667P1423 (from clone DKFZp667P1423) /gb=AL832809 /gi=21733392 /ug=Hs.288997 /len=3567	AL832809	Hs.288997
11061	0.047	clone IMAGE:3887266, mRNA /gb=BC015512 /gi=15930151 /ug=Hs.88013 /len=1505	BC015512	Hs.88013
11062	0.047	F-box only protein 29 (FBXO29), mRNA /cds=(88,1884) /gb=NM_153348 /gi=24158491 /ug=Hs.350985 /len=4874	NM_153348	Hs.350985
11063	9.67E-04	clone alpha_est218/52C1 mRNA sequence /gb=AF001542 /gi=2529714 /ug=Hs.356442 /len=2992	AF001542	Hs.356442
11067	0.00709	mRNA; cDNA DKFZp686H0819 (from clone DKFZp686H0819)	AL832480	
11073	0.023613	cDNA FLJ11174 fis, clone PLACE1007367. /gb=AK002036 /gi=7023674 /ug=Hs.24359 /len=2285	AK002036	Hs.24359
11112	0.006705	K-EST0052858 S14K402 cDNA clone S14K402-16-B02 5', mRNA sequence /clone=S14K402-16-B02 /clone_end=5' /gb=BM769588 /gi=19099203 /ug=Hs.131020 /len=562	BM769588	Hs.131020
11142	0.024176	cDNA: FLJ22447 fis, clone HRC09479. /gb=AK026100 /gi=10438841 /ug=Hs.344000 /len=1659	AK026100	Hs.344000
11181	0.005043	BAC clone RP11-812M14 from 2, complete sequence	AC115115	
11215	0.009316	no match		
11221	0.019043	chromosome 14 DNA sequence BAC C-2062F14 of library CalTech-D from chromosome 14 of complete sequence	AL845552	
11236	0.041391	prenylcysteine lyase, mRNA (cDNA clone MGC:60307 IMAGE:5773815), complete cds (=AB020715.1)	BC051891	
11277	0.031823	ligase I, DNA, ATP-dependent (LIG1), mRNA /cds=(121,2880) /gb=NM_000234 /gi=4557718 /ug=Hs.1770 /len=3083	NM_000234	Hs.1770
11290	0.004758	T-cell leukemia translocation altered gene (TCTA), mRNA /cds=(222,533) /gb=NM_022171 /gi=11560140 /ug=Hs.250894 /len=2146	NM_022171	Hs.250894
11356	0.006705	Mus musculus N-myc downstream regulated 4 (Ndr4), mRNA	NM_145602	Mm.29846
11395	0.029076	LAG1 longevity assurance 2 (S. cerevisiae) (LASS2), mRNA /cds=(50,742) /gb=NM_013384 /gi=9937997 /ug=Hs.285976 /len=1646	NM_013384	Hs.285976
11414	0.023066	solute carrier family 25 (mitochondrial carrier; phosphate carrier), member 3 (SLC25A3), nuclear gene encoding mitochondrial protein, transcript variant 1b, mRNA /cds=(49,1134) /gb=NM_002635 /gi=4505774 /ug=Hs.78713 /len=1330	NM_002635	Hs.78713

11425	0.031823	mucosa associated lymphoid tissue lymphoma translocation gene 1 (MALT1), transcript variant 1, mRNA /cds=(259,2733) /gb=NM_006785 /gi=27886564 /ug=Hs.180566 /len=5029	NM_006785	Hs.180566
11427	0.034783	DKFZP586A0522 protein (DKFZP586A0522), mRNA /cds=(21,755) /gb=NM_014033 /gi=13378140 /ug=Hs.288771 /len=1705	NM_014033	Hs.288771
11434	0.021999	hypothetical protein FLJ20375 (FLJ20375), mRNA /cds=(364,5769) /gb=NM_017794 /gi=23510242 /ug=Hs.274251 /len=6117	NM_017794	Hs.274251
11435	1.54E-05	SR rich protein (DKFZp564B0769), mRNA /cds=(33,2450) /gb=NM_032870 /gi=18699723 /ug=Hs.18368 /len=2663	NM_032870	Hs.18368
11438	0.029076	hypothetical protein FLJ20360 (FLJ20360), mRNA /cds=(80,2305) /gb=NM_017782 /gi=8923334 /ug=Hs.26434 /len=3041	NM_017782	Hs.26434
11455	0.01344	clone IMAGE:3912592, mRNA	BC016846	Hs.334875
11462	0.002767	ta10c07.x1 Soares_total_fetus_Nb2HF8_9w cDNA clone IMAGE:2043660 3', mRNA sequence /clone=IMAGE:2043660 /clone_end=3' /gb=AI581285 /gi=4565661 /ug=Hs.309697 /len=467	AI581285	Hs.309697
11478	0.005343	major histocompatibility complex, class II, DR beta 5 (HLA-DRB5), mRNA /cds=(6,806) /gb=NM_002125 /gi=26665892 /ug=Hs.352392 /len=1171	NM_002125	Hs.352392
11481	0.049002	eps8 binding protein e3B1 mRNA, complete cds	AF006516	Hs.24752
11490	0.045065	enhancer of filamentation 1 (cas-like docking; Crk-associated substrate related) (HEF1), mRNA /cds=(164,2668) /gb=NM_006403 /gi=5453679 /ug=Hs.80261 /len=3817	NM_006403	Hs.80261
11504	0.019991	ankyrin-like with transmembrane domains 1 (ANKTM1), mRNA /cds=(175,3534) /gb=NM_007332 /gi=6601589 /ug=Hs.137674 /len=5190	NM_007332	Hs.137674
11514	0.029076	chromosome 21 open reading frame 33 (C21orf33), mRNA /cds=(85,891) /gb=NM_004649 /gi=5031690 /ug=Hs.182423 /len=1652	NM_004649	Hs.182423
11517	0.012127	fasciculation and elongation protein zeta 1 (zygin I) (FEZ1), transcript variant 1, mRNA	NM_005103	Hs.79226
11537	8.41E-04	UI-H-DF1-auf-c-04-0-UI.s1 NCI_CGAP_DF1 cDNA clone IMAGE:5868603 3', mRNA sequence /clone=IMAGE:5868603 /clone_end=3' /gb=BM992029 /gi=19711418 /ug=Hs.358825 /len=1052	BM992029	Hs.358825
11549	0.018141	DNA sequence from clone RP5-975D15 on chromosome 1p31.3-32.2, complete sequence	AL136120	
11582	0.002292	RC1-NN0073-090500-012-f02 NN0073 cDNA, mRNA sequence /gb=AW898615 /gi=8062820 /ug=Hs.130729 /len=660	AW898615	Hs.130729
11587	0.01564	no match		
11590	0.031823	UI-E-EJ0-aig-j-08-0-UI.s1 UI-E-EJ0 cDNA clone UI-E-EJ0-aig-j-08-0-UI 3', mRNA sequence /clone=UI-E-EJ0-aig-j-08-0-UI /clone_end=3' /gb=BM682503 /gi=18992399 /ug=Hs.446242 /len=1052	BM682503	Hs.446242



11630	0.002292	uncharacterized hypothalamus protein HT007 (HT007), mRNA /cds=(228,887) /gb=NM_018480 /gi=8923801 /ug=Hs.24371 /len=1172	NM_018480	Hs.24371
11681	0.045065	chromosome 8, clone CTD-3214K23, complete sequence	AC113139	
11711	0.030424	EH-domain containing 1 (EHD1), mRNA /cds=(247,1851) /gb=NM_006795 /gi=5803008 /ug=Hs.155119 /len=3508	NM_006795	Hs.155119
11725	0.02533	hypothetical protein FLJ13657 (FLJ13657), mRNA /cds=(88,1173) /gb=NM_024828 /gi=13376229 /ug=Hs.178357 /len=2252	NM_024828	Hs.178357
11729	0.014875	hyaluronoglucosaminidase 3 (HYAL3), mRNA /cds=(274,1527) /gb=NM_003549 /gi=15208650 /ug=Hs.129910 /len=1942	NM_003549	Hs.129910
11743	0.036347	CD74 antigen (invariant polypeptide of major histocompatibility complex, class II antigen-associated) (CD74), mRNA /cds=(8,706) /gb=NM_004355 /gi=10835070 /ug=Hs.84298 /len=1304	NM_004355	Hs.84298
11782	0.012769	apoptosis-related protein PNAS-1 (LOC51275), mRNA /cds=(543,767) /gb=NM_016534 /gi=7706078 /ug=Hs.334467 /len=823	NM_016534	Hs.334467
11854	0.003986	partial RANBP7 gene for RanBP7/importin7 and partial ZNF143 gene	AJ295844	
11890	0.010925	similar to RIKEN cDNA 5730528L13 gene (MGC17337), mRNA /cds=(68,895) /gb=NM_080655 /gi=18087818 /ug=Hs.78531 /len=1175	NM_080655	Hs.78531
11898	0.02533	intersectin 2 (ITSN2), transcript variant 1, mRNA /cds=(242,5332) /gb=NM_006277 /gi=22325384 /ug=Hs.166184 /len=6092	NM_006277	Hs.166184
11899	0.020975	hypothetical protein LOC51244 (LOC51244), mRNA /cds=(340,1233) /gb=NM_016474 /gi=24475969 /ug=Hs.158006 /len=1614	NM_016474	Hs.158006
11941	0.00136	uncharacterized hematopoietic stem/progenitor cells protein MDS031 (MDS031), mRNA /cds=(35,532) /gb=NM_018466 /gi=20070304 /ug=Hs.110853 /len=1358	NM_018466	Hs.110853
11943	0.020975	chromosome 10 open reading frame 4 (C10orf4), mRNA /cds=(199,1146) /gb=NM_145246 /gi=24432066 /ug=Hs.351929 /len=2400	NM_145246	Hs.351929
11951	0.041391	chromosome 17, clone RP11-567L7, complete sequence	AC091059	
11954	0.021999	Mus musculus apoptosis antagonizing transcription factor (Aatf), mRNA	NM_019816	Mm.18994
11973	0.031823	mitochondrion, complete genome	NC_001807	
11983	0.047	RAS-like, estrogen-regulated, growth-inhibitor (RERG), mRNA /cds=(291,890) /gb=NM_032918 /gi=14249703 /ug=Hs.21594 /len=2240	NM_032918	Hs.21594
12004	0.020975	chromosome 4 clone RP11-442P12, complete sequence	AC005798	

12049	4.72E-04	TAF2 RNA polymerase II, TATA box binding protein (TBP)-associated factor, 150kDa (TAF2), mRNA /cds=(271,3870) /gb=NM_003184 /gi=20357590 /ug=Hs.122752 /len=5019	NM_003184	Hs.122752
12061	0.033276	cDNA FLJ36838 fis, clone ASTRO2011426. /gb=AK094157 /gi=21753159 /ug=Hs.407030 /len=2646	AK094157	Hs.407030
12081	0.031823	hypothetical protein FLJ13855 (FLJ13855), mRNA /cds=(328,1068) /gb=NM_023079 /gi=20149671 /ug=Hs.168232 /len=3053	NM_023079	Hs.168232
12088	0.039649	cDNA FLJ37791 fis, clone BRHIP3000131. /gb=AK095110 /gi=21754304 /ug=Hs.350534 /len=3820	AK095110	Hs.350534
12098	0.043196	BAC clone RP11-722P9 from 2, complete sequence	AC133534	
12120	0.019991	mRNA; cDNA DKFZp761O0611 (from clone DKFZp761O0611) /gb=AL834155 /gi=21739631 /ug=Hs.22969 /len=4502	AL834155	Hs.22969
12126	0.023066	clone IMAGE:5273893, mRNA	BC041001	Hs.7194
12160	0.016439	DKFZp564P1871_s1 564 (synonym: hfbr2) cDNA clone DKFZp564P1871 3', mRNA sequence /clone=DKFZp564P1871 /clone_end=3' /gb=AL037446 /gi=5406837 /ug=Hs.208747 /len=556	AL037446	Hs.208747
12166	0.00599	ot86d06.s1 Soares_total_fetus_Nb2HF8_9w cDNA clone IMAGE:1623659 3', mRNA sequence /clone=IMAGE:1623659 /clone_end=3' /gb=AI016364 /gi=3230700 /ug=Hs.131059 /len=478	AI016364	Hs.131059
12202	0.037968	F-box and leucine-rich repeat protein 3A (FBXL3A), mRNA /cds=(298,1584) /gb=NM_012158 /gi=16306583 /ug=Hs.7540 /len=3489	NM_012158	Hs.7540
12232	0.047	protein disulfide isomerase related protein (calcium-binding protein, intestinal-related) (ERP70), mRNA /cds=(243,2180) /gb=NM_004911 /gi=21624646 /ug=Hs.93659 /len=2930	NM_004911	Hs.93659
12261	0.041391	UI-E-EO1-aid-o-06-0-UI.s1 UI-E-EO1 cDNA clone UI-E-EO1-aid-o-06-0-UI 3', mRNA sequence /clone=UI-E-EO1-aid-o-06-0-UI /clone_end=3' /gb=BM677516 /gi=18987412 /ug=Hs.443680 /len=1044	BM677516	Hs.443680
12272	0.01344	chromosome 5 clone CTB-43D14, complete sequence	AC022096	
12329	0.036347	DNA sequence from clone RP11-438P12 on chromosome 10, complete sequence	AL157397	
12360	0.02533	chromosome 3 clone RP11-322K13, complete sequence	AC113169	
12374	0.005343	clone IMAGE:5223469, mRNA	BC043380	Hs.221776
12384	0.029076	cDNA FLJ31039 fis, clone HSYRA2000221	AK055601	Hs.311977
12408	0.049002	phosphatidic acid phosphatase type 2B (PPAP2B), transcript variant 2, mRNA	NM_177414	
12447	0.037968	cDNA FLJ12049 fis, clone HEMBB1001996. /gb=AK022111 /gi=10433434 /ug=Hs.171395 /len=2100	AK022111	Hs.171395
12467	0.02533	chromosome 15, clone RP11-26F2, complete sequence	AC011767	
12481	0.026531	BAC clone RP11-738E22 from 4, complete sequence	AC069307	

12483	0.036347	mRNA; cDNA DKFZp667M218 (from clone DKFZp667M218) /gb=AL713739 /gi=19584478 /ug=Hs.375787 /len=2989	AL713739	Hs.375787
12490	0.02533	similar to hypothetical protein LOC283596 (LOC338929), mRNA	XM_211115	
12503	0.045065	AV738079 CB cDNA clone CBFDPF09 5', mRNA sequence /clone=CBFDPF09 /clone_end=5' /gb=AV738079 /gi=10855660 /ug=Hs.378390 /len=238	AV738079	Hs.378390
12539	0.001272	UI-E-CK1-afh-a-18-0-UI.r1 UI-E-CK1 cDNA clone UI-E-CK1-afh-a-18-0-UI 5', mRNA sequence /clone=UI-E-CK1-afh-a-18-0-UI /clone_end=5' /gb=BM702618 /gi=19015876 /ug=Hs.103381 /len=1069	BM702618	Hs.103381
12546	0.008362	cDNA FLJ38331 fis, clone FCBBF3025285, moderately similar to Mus musculus peripheral benzodiazepine receptor associated protein (Pap7) mRNA. /gb=AK095650 /gi=21754954 /ug=Hs.9052 /len=3547	AK095650	Hs.9052
12554	0.043196	mRNA; cDNA DKFZp667D2123 (from clone DKFZp667D2123) /gb=AL832786 /gi=21733368 /ug=Hs.283643 /len=3000	AL832786	Hs.283643
12555	0.047	DNA sequence from clone RP1-84N20 on chromosome 6 Contains a novel gene, ESTs, STSs, GSSs and two CpG islands, complete sequence	AL136128	
12653	0.034783	cDNA FLJ10247 fis, clone HEMBB1000705. /gb=AK001109 /gi=7022167 /ug=Hs.24610 /len=1601	AK001109	Hs.24610
12673	0.049002	cDNA FLJ37017 fis, clone BRACE2010642. /gb=AK094336 /gi=21753377 /ug=Hs.27280 /len=2160	AK094336	Hs.27280
12683	1.34E-04	yi11a11.s1 Soares placenta Nb2HP cDNA clone IMAGE:138908 3', mRNA sequence /clone=IMAGE:138908 /clone_end=3' /gb=R62551 /gi=834430 /ug=Hs.143796 /len=344	R62551	Hs.143796
12687	0.027779	wr41b04.x1 NCI_CGAP_Pr28 cDNA clone IMAGE:2490223 3', mRNA sequence /clone=IMAGE:2490223 /clone_end=3' /gb=AI972618 /gi=5769444 /ug=Hs.370369 /len=225	AI972618	Hs.370369
12697	0.029076	mitochondrion, complete genome	NC_001807	
12700	0.006705	NADH:ubiquinone oxidoreductase MLRQ subunit (LOC56901), mRNA /cds=(274,537) /gb=NM_020142 /gi=20127589 /ug=Hs.110024 /len=1284	NM_020142	Hs.110024
12713	0.041391	DNA sequence from clone RP11-248G5 on chromosome 13q14.3-21.1, complete sequence	AL139082	
12757	0.002017	chromosome 10 clone RP11-348G8, complete sequence	AC073160	
12759	0.039649	UI-H-BI1-aee-g-08-0-UI.s1 NCI_CGAP_Sub3 cDNA clone IMAGE:2719286 3', mRNA sequence /clone=IMAGE:2719286 /clone_end=3' /gb=AW139957 /gi=6144675 /ug=Hs.396536 /len=744	AW139957	Hs.396536

12787	0.030424	similar to RIKEN cDNA 1110018M03, clone MGC:24932 IMAGE:4938507, mRNA, complete cds /cds=(218,853) /gb=BC026873 /gi=20073062 /ug=Hs.32478 /len=1826	BC026873	Hs.32478
12789	0.037968	Similar to PP4189, clone MGC:45566 IMAGE:4385905, mRNA, complete cds	BC035604	Hs.170453
12803	0.020975	Mus musculus adult male corpora quadrigemina cDNA, RIKEN full-length enriched library, clone:B230344L23 product:protocadherin 10, full insert sequence	AK046143	Mm.44368
12855	0.018141	UI-E-EJ0-aik-i-20-0-UI.r1 UI-E-EJ0 cDNA clone UI-E-EJ0-aik-i-20-0-UI 5', mRNA sequence /clone=UI-E-EJ0-aik-i-20-0-UI /clone_end=5' /gb=BM727413 /gi=19048746 /ug=Hs.112619 /len=1667	BM727413	Hs.112619
12856	0.014875	cDNA FLJ33798 fis, clone CTONG2000063. /gb=AK091117 /gi=21749410 /ug=Hs.7921 /len=3551	AK091117	Hs.7921
12870	0.024176	chromosome 5 clone CTB-113P19, complete sequence	AC011374	
12896	0.02533	cDNA FLJ30298 fis, clone BRACE2003172. /gb=AK054860 /gi=16549479 /ug=Hs.351546 /len=2659	AK054860	Hs.351546
12903	0.037968	cDNA FLJ33097 fis, clone TRACH2000775. /gb=AK057659 /gi=16553423 /ug=Hs.415317 /len=2977	AK057659	Hs.415317
12933	0.019047	cDNA FLJ34764 fis, clone NT2NE2002311. /gb=AK092083 /gi=21750590 /ug=Hs.111583 /len=2552	AK092083	Hs.111583
12934	0.014142	cDNA FLJ14193 fis, clone NT2RP3001115. /gb=AK024255 /gi=10436585 /ug=Hs.162990 /len=2308	AK024255	Hs.162990
12940	0.033276	cDNA FLJ31626 fis, clone NT2RI2003317. /gb=AK056188 /gi=16551523 /ug=Hs.375198 /len=2041	AK056188	Hs.375198
12941	0.014875	MR2-CI0186-291100-010-a06 CI0186 cDNA, mRNA sequence /gb=BF814502 /gi=12147047 /ug=Hs.446594 /len=530	BF814502	Hs.446594
12960	0.049002	hypothetical protein FLJ25157 (FLJ25157), mRNA /cds=(157,762) /gb=NM_152653 /gi=22749326 /ug=Hs.108323 /len=1541	NM_152653	Hs.108323
13010	0.043196	cDNA FLJ32247 fis, clone PROST1000120. /gb=AK056809 /gi=16552317 /ug=Hs.293663 /len=3019	AK056809	Hs.293663
13014	0.012127	cDNA FLJ13334 fis, clone OVARC1001846. /gb=AK023396 /gi=10435315 /ug=Hs.269091 /len=2361	AK023396	Hs.269091
13026	0.033276	blank		
13039	0.012769	601571978T1 NIH_MGC_55 cDNA clone IMAGE:3838987 3', mRNA sequence /clone=IMAGE:3838987 /clone_end=3' /gb=BE748785 /gi=10162777 /ug=Hs.334633 /len=801	BE748785	Hs.334633
13067	0.039649	ARF 9		

13081	0.024176	12 BAC RP11-302B13 (Roswell Park Cancer Institute BAC Library) complete sequence	AC073610	
13106	0.030424	growth arrest and DNA-damage-inducible, beta (GADD45B), mRNA /cds=(101,586) /gb=NM_015675 /gi=9945331 /ug=Hs.110571 /len=1121	NM_015675	Hs.110571
13130	0.009828	mRNA for KIAA0622 protein, partial cds. /cds=(1,3870) /gb=AB014522 /gi=3327057 /ug=Hs.11238 /len=6951	AB014522	Hs.11238
13136	0.047	mRNA for KIAA0515 protein, partial cds. /cds=(1,2014) /gb=AB011087 /gi=3043553 /ug=Hs.405891 /len=6335	AB011087	Hs.405891
13163	0.006339	RNA polymerase I subunit (RPA40), mRNA /cds=(23,1051) /gb=NM_004875 /gi=4759045 /ug=Hs.5409 /len=1103	NM_004875	Hs.5409
13181	0.005658	inositol(myo)-1(or 4)-monophosphatase 2 (IMPA2), mRNA /cds=(215,1081) /gb=NM_014214 /gi=7657235 /ug=Hs.5753 /len=1500	NM_014214	Hs.5753
13185	0.030424	mRNA; cDNA DKFZp667H216 (from clone DKFZp667H216) /gb=AL833204 /gi=21733834 /ug=Hs.356145 /len=3782	AL833204	Hs.356145
13202	0.009316	hypothetical protein FLJ12806 (FLJ12806), mRNA /cds=(158,1078) /gb=NM_022831 /gi=12383075 /ug=Hs.107637 /len=2485	NM_022831	Hs.107637
13221	0.027779	UI-H-DF1-auf-c-04-0-UI.s1 NCI_CGAP_DF1 cDNA clone IMAGE:5868603 3', mRNA sequence /clone=IMAGE:5868603 /clone_end=3' /gb=BM992029 /gi=19711418 /ug=Hs.358825 /len=1052	BM992029	Hs.358825
13283	0.005343	SEC24 related gene family, member D (S. cerevisiae) (SEC24D), mRNA /cds=(201,3299) /gb=NM_014822 /gi=7662658 /ug=Hs.19822 /len=3988	NM_014822	Hs.19822
13284	0.031823	mRNA full length insert cDNA clone EUROIMAGE 1709050	AJ420494	Hs.348999
13294	0.020975	actin related protein 2/3 complex, subunit 1A, 41kDa (ARPC1A), mRNA /cds=(148,1260) /gb=NM_006409 /gi=22907051 /ug=Hs.90370 /len=1619	NM_006409	Hs.90370
13341	0.034783	leucine aminopeptidase 3 (LAP3), mRNA /cds=(187,1746) /gb=NM_015907 /gi=7705687 /ug=Hs.182579 /len=2147	NM_015907	Hs.182579
13342	0.01344	RNA guanylyltransferase and 5'-phosphatase (RNGTT), mRNA /cds=(289,2082) /gb=NM_003800 /gi=4506562 /ug=Hs.27345 /len=4546	NM_003800	Hs.27345
13348	0.036347	hypothetical protein FLJ14153 (FLJ14153), mRNA /cds=(31,1428) /gb=NM_022736 /gi=12232392 /ug=Hs.7503 /len=2161	NM_022736	Hs.7503
13351	0.030424	basic transcription factor 3 (BTF3), mRNA /cds=(240,728) /gb=NM_001207 /gi=20070129 /ug=Hs.101025 /len=952	NM_001207	Hs.101025
13353	0.012769	transcription termination factor-like protein (LOC80298), mRNA /cds=(341,1498) /gb=NM_025198 /gi=21314735 /ug=Hs.5009 /len=1792	NM_025198	Hs.5009

13354	0.012769	catenin, beta like 1 (CTNBL1), mRNA /cds=(95,1786) /gb=NM_030877 /gi=18644733 /ug=Hs.178576 /len=1900	NM_030877	Hs.178576
13362	0.041391	cDNA FLJ11469 fis, clone HEMBA1001658. /gb=AK021531 /gi=10432731 /ug=Hs.224398 /len=1665	AK021531	Hs.224398
13368	0.034783	FGF receptor activating protein 1 (FRAG1), mRNA /cds=(129,1076) /gb=NM_014489 /gi=7657101 /ug=Hs.133968 /len=2040	NM_014489	Hs.133968
13369	0.012127	growth hormone inducible transmembrane protein (GHITM), mRNA /cds=(130,1089) /gb=NM_014394 /gi=7657479 /ug=Hs.433957 /len=2374	NM_014394	Hs.433957
13377	0.039649	hypothetical protein FLJ22729 (FLJ22729), mRNA /cds=(603,1079) /gb=NM_024683 /gi=13375953 /ug=Hs.94891 /len=1278	NM_024683	Hs.94891
13379	0.043196	patched related protein translocated in renal cancer (TRC8), mRNA /cds=(215,2209) /gb=NM_007218 /gi=21314653 /ug=Hs.28285 /len=2481	NM_007218	Hs.28285
13389	0.034783	brain protein 44-like (BRP44L), mRNA /cds=(123,452) /gb=NM_016098 /gi=7706368 /ug=Hs.108725 /len=988	NM_016098	Hs.108725
13408	0.006705	no match		
13417	0.005658	BAC clone RP11-208G20 from 7, complete sequence	AC104692	
13427	0.02533	DNA sequence from clone RP11-164D15 on chromosome 9, complete sequence	AL590642	
13476	0.031823	chromosome 5 clone CTC-484M2, complete sequence	AC020899	
13478	0.045065	cDNA FLJ33928 fis, clone CTONG2017444	AK091247	Hs.16603
13491	0.003535	hypothetical protein MGC33602 (MGC33602), mRNA /cds=(140,748) /gb=NM_152391 /gi=22748836 /ug=Hs.274415 /len=1790	NM_152391	Hs.274415
13510	0.049002	uncharacterized hematopoietic stem/progenitor cells protein MDS031 (MDS031), mRNA /cds=(35,532) /gb=NM_018466 /gi=20070304 /ug=Hs.110853 /len=1358	NM_018466	Hs.110853
13513	0.008362	hypothetical protein FLJ20203 (FLJ20203), mRNA	NM_032292	Hs.20594
13552	0.009828	cDNA FLJ36224 fis, clone THYMU2000990. /gb=AK093543 /gi=21752443 /ug=Hs.250367 /len=2299	AK093543	Hs.250367
13554	0.012127	cyclin M3 (CNNM3), mRNA /cds=(99,1247) /gb=NM_017623 /gi=20127562 /ug=Hs.44095 /len=2234	NM_017623	Hs.44095
13582	0.009828	mRNA; cDNA DKFZp586M1819 (from clone DKFZp586M1819) /cds=(1,795) /gb=AL834255 /gi=21739805 /ug=Hs.355753 /len=1723	AL834255	Hs.355753
13593	0.011512	low density lipoprotein-related protein-associated protein 1 (alpha-2-macroglobulin receptor-associated protein 1) (LRPAP1), mRNA /cds=(14,1087) /gb=NM_002337 /gi=4505020 /ug=Hs.75140 /len=1493	NM_002337	Hs.75140

13605	0.00423	unr-interacting protein (UNRIP), mRNA /cds=(297,1349) /gb=NM_007178 /gi=20149591 /ug=Hs.3727 /len=1867	NM_007178	Hs.3727
13654	0.039649	hypothetical protein LOC92597 (LOC92597), mRNA /cds=(151,801) /gb=NM_173468 /gi=27735028 /ug=Hs.31422 /len=6956	NM_173468	Hs.31422
13655	3.49E-04	vacuolar protein sorting 29 (yeast) (VPS29) transcript variant 2, mRNA /cds=(61,621) /gb=NM_057180 /gi=17402911 /ug=Hs.69192 /len=1107	NM_057180	Hs.69192
13675	0.023066	guanine nucleotide binding protein beta subunit 4 (GNB4), mRNA /cds=(281,1303) /gb=NM_021629 /gi=20357531 /ug=Hs.172654 /len=3302	NM_021629	Hs.172654
13691	0.023066	POP4 (processing of precursor, S. cerevisiae) (POP4), mRNA /cds=(26,688) /gb=NM_006627 /gi=5729985 /ug=Hs.421667 /len=1133	NM_006627	Hs.421667
13692	0.019991	cDNA FLJ38575 fis, clone HCHON2007046. /gb=AK095894 /gi=21755244 /ug=Hs.376206 /len=2134	AK095894	Hs.376206
13695	0.019991	BAC clone RP11-1110 from 4, complete sequence	AC096712	
13698	0.010925	hypothetical protein DKFZp564F013 (DKFZP564F013), mRNA /cds=(107,2194) /gb=NM_020432 /gi=24308192 /ug=Hs.128653 /len=4572	NM_020432	Hs.128653
13700	0.01564	cDNA FLJ13127 fis, clone NT2RP3002911	AK023189	Hs.143917
13701	0.01564	v-raf murine sarcoma 3611 viral oncogene 1 (ARAF1), mRNA /cds=(203,2023) /gb=NM_001654 /gi=4502192 /ug=Hs.77183 /len=2466	NM_001654	Hs.77183
13703	0.027779	cDNA FLJ35748 fis, clone TESTI2004459, highly similar to cholinephosphotransferase 1 alpha mRNA	AK093067	Hs.171889
13767	0.045065	AGENCOURT_8928839 NIH_MGC_142 cDNA clone IMAGE:6498777 5', mRNA sequence /clone=IMAGE:6498777 /clone_end=5' /gb=BU506761 /gi=22812994 /ug=Hs.355839 /len=1148	BU506761	Hs.355839
13778	0.011512	clone IMAGE:5287569, mRNA	BC047058	
13859	0.008828	Repetitive Sequence		
13946	0.041391	cDNA FLJ13536 fis, clone PLACE1006521. /gb=AK023598 /gi=10435577 /ug=Hs.11493 /len=2132	AK023598	Hs.11493
13979	0.041391	no match		
13983	0.049002	genomic DNA, chromosome 8q23, clone: KB2047A4	AP003477	
13985	0.005043	3 BAC RP11-115H18 (Roswell Park Cancer Institute BAC Library) complete sequence	AC068765	
13992	0.021999	DNA sequence from clone RP4-657E11 on chromosome 1p35.1-36.23, complete sequence	AL035413	
14011	0.036347	tm42d10.x1 NCI_CGAP_Kid11 cDNA clone IMAGE:2160787 3', mRNA sequence /clone=IMAGE:2160787 /clone_end=3' /gb=AI498708 /gi=4390690 /ug=Hs.170849 /len=453	AI498708	Hs.170849

14042	7.84E-04	UI-H-DT0-atw-d-24-0-UI.s1 NCI_CGAP_DT0 cDNA clone IMAGE:5865191 3', mRNA sequence /clone=IMAGE:5865191 /clone_end=3' /gb=BM994000 /gi=19718901 /ug=Hs.395775 /len=1181	BM994000	Hs.395775
14048	0.001454	glutamyl-prolyl-tRNA synthetase (EPRS), mRNA /cds=(59,4381) /gb=NM_004446 /gi=4758293 /ug=Hs.55921 /len=4586	NM_004446	Hs.55921
14057	0.037968	cDNA FLJ90553 fis, clone OVARC1000853. /cds=(116,748) /gb=AK075034 /gi=22760867 /ug=Hs.406158 /len=1673	AK075034	Hs.406158
14071	0.037968	genomic DNA, chromosome 21q22.2, DSCR region, clone D47-S479, segment 15/16, complete sequence	AP000163	
14097	0.01564	MR0-HT0559-290500-027-d10 HT0559 cDNA, mRNA sequence /gb=BE708268 /gi=10096533 /ug=Hs.209224 /len=619	BE708268	Hs.209224
14109	0.026531	wq35e02.x1 NCI_CGAP_GC6 cDNA clone IMAGE:2473274 3', mRNA sequence /clone=IMAGE:2473274 /clone_end=3' /gb=AI950442 /gi=5742752 /ug=Hs.176956 /len=496	AI950442	Hs.176956
14137	0.031823	ribosomal protein S4, X-linked (RPS4X), mRNA /cds=(36,827) /gb=NM_001007 /gi=17981705 /ug=Hs.389933 /len=916	NM_001007	Hs.389933
14145	5.88E-04	mitochondrion, complete genome	NC_001807	
14160	0.039649	BAC clone RP11-703L16 from 2, complete sequence	AC068039	
14179	0.018141	BX116503 Soares_multiple_sclerosis_2NbHMSP cDNA clone IMAGp998O04618, mRNA sequence /clone=IMAGp998O04618; IMAGE:279771 /gb=BX116503 /gi=27880522 /ug=Hs.247117 /len=594	BX116503	Hs.247117
14193	0.049002	hypothetical protein FLJ14494 (FLJ14494), mRNA /cds=(9,1142) /gb=NM_032795 /gi=14249469 /ug=Hs.322406 /len=2146	NM_032795	Hs.322406
14226	0.024176	AGENCOURT_6480263 NIH_MGC_92 cDNA clone IMAGE:5575699 5', mRNA sequence /clone=IMAGE:5575699 /clone_end=5' /gb=BM475617 /gi=18524659 /ug=Hs.445483 /len=1135	BM475617	Hs.445483
14267	0.008362	yo08g06.r1 Soares adult brain N2b5HB55Y cDNA clone IMAGE:177370 5', mRNA sequence /clone=IMAGE:177370 /clone_end=5' /gb=H40746 /gi=916798 /ug=Hs.408835 /len=500	H40746	Hs.408835
14286	0.014142	mRNA; cDNA DKFZp434J214 (from clone DKFZp434J214); partial cds /cds=(1,1082) /gb=AL080156 /gi=5262614 /ug=Hs.12813 /len=2749	AL080156	Hs.12813
14297	0.036347	cDNA FLJ34184 fis, clone FCBBF3017024. /gb=AK091503 /gi=21749887 /ug=Hs.146217 /len=2896	AK091503	Hs.146217
14382	0.034783	mRNA; cDNA DKFZp434P1018 (from clone DKFZp434P1018); partial cds	AL137527	Hs.289038
14399	0.014141	mRNA; cDNA DKFZp451E0516 (from clone DKFZp451E0516)	AL832578	Hs.306219



14426	0.021999	AV700508 GKC cDNA clone GKCGLC05 3', mRNA sequence /clone=GKCGLC05 /clone_end=3' /gb=AV700508 /gi=10302479 /ug=Hs.285895 /len=559	AV700508	Hs.285895
14444	0.017272	mitochondrion, complete genome	NC_001807	
14450	0.002292	no match		
14485	0.036347	no match		
14488	0.030424	no match		
14520	0.007917	cDNA FLJ90504 fis, clone NT2RP3004090, weakly similar to GOLIATH PROTEIN. /cds=(103,1305) /gb=AK074985 /gi=22760786 /ug=Hs.171802 /len=2452	AK074985	Hs.171802
14527	0.004487	URB mRNA, complete cds /cds=(146,2998) /gb=AF506819 /gi=21039408 /ug=Hs.356289 /len=3320	AF506819	Hs.356289
14556	0.017272	hypothetical protein FLJ23221 (FLJ23221), mRNA /cds=(24,419) /gb=NM_024579 /gi=13375757 /ug=Hs.18397 /len=519	NM_024579	Hs.18397
14558	0.041391	12 BAC RP11-463O12 (Roswell Park Cancer Institute BAC Library) complete sequence	AC027290	
14586	0.043193	clone IMAGE:125405, mRNA sequence /gb=AF339770 /gi=13507308 /ug=Hs.221635 /len=790	AF339770	Hs.221635
14631	0.037968	DNA sequence from clone RP11-434B7 on chromosome 1, complete sequence	AL583826	
14721	0.018141	UI-H-ED0-axn-i-09-0-UI.s1 NCI_CGAP_ED0 cDNA clone UI-H-ED0-axn-i-09-0-UI 3', mRNA sequence /clone=UI-H-ED0-axn-i-09-0-UI /clone_end=3' /gb=CA445401 /gi=24809821 /ug=Hs.204930 /len=725	CA445401	Hs.204930
14772	0.029076	AJ318805 adipose tissue cDNA clone 2040, mRNA sequence /clone=2040 /gb=AJ318805 /gi=18141682 /ug=Hs.86538 /len=5223	AJ318805	Hs.86538
14775	0.027779	Chromosome X PAC RPC11-290C9 from the Pieter de Jong PAC library; complete sequence	AC002404	
14828	0.034783	similar to squamous cell carcinoma-related protein-1 (LOC284248), mRNA	XM_209090	
14858	0.034783	chromosome 5 clone RP11-175M2, complete sequence	AC016632	
14882	0.029076	DKFZp434C022_s1 434 (synonym: htes3) cDNA clone DKFZp434C022 3', mRNA sequence /clone=DKFZp434C022 /clone_end=3' /gb=AL044366 /gi=5432588 /ug=Hs.165805 /len=668	AL044366	Hs.165805
14884	0.030424	602043661F1 NCI_CGAP_Brn67 cDNA clone IMAGE:4181462 5', mRNA sequence /clone=IMAGE:4181462 /clone_end=5' /gb=BF528488 /gi=11615851 /ug=Hs.433462 /len=885	BF528488	Hs.433462
14924	0.014875	no match		
14954	0.014141	DNA sequence from clone RP4-621O15 on chromosome 1, complete sequence	AL713899	
14967	0.036347	no match		